

# STIC Search Report Biotech-Chem Library

STIC Database Tracking Number

TO: Emily M Le

**Location: 3c35/3c18** 

Art Unit: 1648

Monday, May 16, 2005

Case Serial Number: 08/869386

From: Noble Jarrell

**Location: Biotech-Chem Library** 

**Rem 1B71** 

Phone: 272-2556

Noble.jarrell@uspto.gov

#### Search Notes

### **Protein Sequence Searches - February 2005**

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.



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### 153406 CRFE

#### Jarrell, Noble

From:

Le, Emily

Sent:

Monday, May 16, 2005 12:04 PM

To:

Jarrell, Noble

Subject:

FW: Sequence Search: 08869386

#### update:

Please provide a search for the following:

1. RAFVTIGK, which is SEQ ID NO: 5 in the above case.

2. SEQ ID NO: 1

3. SEQ ID NO: 3

Please also limit the size to no more than 25 amino acids.

Thanks, Noble.

#### Emily

----Original Message-----

Le, Emily

Sent:

Friday, May 13, 2005 4:13 PM

To:

Jarrell, Noble

Subject:

Sequence Search: 08869386

#### Noble,

Please provide a search for the following:

1. RAFVTIGK

Please also limit the size to no more than 25 amino acids.

#### Thanks!

Emily Le Office, Rem 3C35 Mailbox, Rem 3C18 Tel., 2-0903

noll

May 162005 3NA, Compugur 10 prep

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Commerical

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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

May 16, 2005, 07:26:50 ; Search time 167 Seconds (without alignments) 24.531 Million cell updates/sec Run on:

1 rafvtigk 8 Title: Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1612378 seqs, 512079187 residues Searched:

16988 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

UniProt\_03:\*
1: uniprot\_sprot:\*
2: uniprot\_trembl:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

SUMMARIES

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Submitted (FEB-2003) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                        MEDLINE-94237431; PubMed-7514146; DOI=10.1016/0378-1097(94)90053-1; Mikkonen M., Vuoristo J., Alatossava T.; "Ribosome binding site consensus sequence of Lactobacillus delbrueckii
              Mikkonen M., Dupont L., Alatossava T., Ritzenthaler P.; "Defective site-specific integration elements are present in the genome of virulent bacteriophage LL-H of Lactobacillus delbrueckii."; Appl. Environ. Microbiol. 62:1847-1851(1996).
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"Complex DNA rearrangements in the att-integration genome regions in related virulent and temperate phages of Lactobacillus delbrueckii.";
Submitted (JUN-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; L42315; AAB06218.1;
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0; Mismatches 1,
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                                                                                                                                                                                                                                                                subsp. lactis bacteriophage LL-H.";
FEMS Microbiol. Lett. 116:315-320(1994)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Fragment)
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01-MAY-2000 (TrEMBLrel. 13, Last seq
MEDLINE=96209253; PubMed=8633887;
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10 AA; 1163 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-2002 (TrEMBLrel. 21, MEPRIN-METALLOENDOPEPTIDASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          56.4%;
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Matches 4; Conserv
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SEQUENCE FROM N.A.
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Gaps

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SEQUENCE FROM N.A.

MEDLINE-29255649, Pubmed=9100996;
Rencher S.D., Lockey T.D., Slobod K.S., Hurwitz J.L.;

"Drift from the GPGRAF HIV-1 envelope V3 crown sequence in a North Anatican liner city.";

AIDS Res. Hum. Retroviruses 13:527-528(1997).

EMBL; U81241; AAB53843.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Characterization of placentation-specific binucleate cell glycoproteins possessing a novel carbohydrate. Evidence for a new family of pregnancy-associated molecules."; family of pregnancy-associated molecules."; placentales. Bates 268:26679-26685 (1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Cetartiodactyla, Ruminantia, Pecora, Bovidae;
Caprinae, Ovis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
PLACENTATION-specific BINUCLEATE cell GLYCOPROTEIN=62 kDa major phytohemaggluthin-binding protein.
Ovis aries (Sheep).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=94075364; PubMed=8253801;
Atkinson Y.H., Gogolin-Ewens K.J., Hounsell E.F., Davies M.J.,
Brandon M.R., Seamark R.F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 22; DB 2; Length 25;
Pred. No. 2.2e+03;
----has 2; Indels
                                                                                                                                                                                         Length 23;
                                                                                                                                                                                                                                     2; Indels
                                         STRAIN=L2 434B;
Wang L., Steenburg S.D., Zheng Y., Larsen S.H.;
Wang Le, Steenburg S.D., the EMBL/GenBank/DDBJ databases.
Submitted (AUG-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF08729; AAD04067.1; -.
                                                                                                                                                 23 AA; 2596 MW; 95DA4A282DC334CE CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25 AA; 2778 MW; 1053D1A02B4BA442 CRC64;
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
                                                                                                                                                                                         Score 22; DB 2;
Pred. No. 2e+03;
1; Mismatches
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                                                                                                                                                                                           56.4%;
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Local Similarity 57.1%;
les 4; Conservative
                                                                                                                                                                                                                                          4; Conservative
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Best Local Similarity
Matches 4; Conserv
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                        FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Caprinae, Ovis.
NCBI_TaxID=9940;
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[1]
SEQUENCE F
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Matches
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                                                                                                                                  Lycopersicon esculentum (Tomato).
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnollophyta, eudicotyledons; core eudicots; asterids;
lamiids; Solanales; Solanaceae; Solanum.
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                                                                                                                                                                                                                                                                                               MEDLINE=21323136; PubMed=11430427; DOI=10.1023/A:1010625203485; Hoeberichts F.A., Orzaez D., van der Plas L.H.W., Woltering E.J.; Changes in gene expression during programmed cell death in tomato cell suspensions."; Paus Mol. 810-1. 45:641-654(2001).

EMBL; AJ250000; CAB61884.1; -.
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01-MAY.2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Hypothetical protonents (Fragment).
Chlamydia trachomatis.
Bacteria: Chlamydiae; Chlamydiae; Chlamydia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
Virulence protein PGP7-D (Fragment).
Chlamydia trachomatis.
Chlamydia trachomatis.
Bacteria, Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia.
NCBI_TaxID=813;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 22; DB 2; Length 17;
Pred, No. 1.5e+03;
1; Mismatches 2; Indels
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57.1%; Pred. No. 1.9e+03;
tive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wang L., Steenburg S.D., Zheng Y., Lareen S.H.;
Submitted (AUG-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF087312; AAD04087.1; -.
Hypothetical protein. 21
                                                                                                                                                                                                                                                                                                                                                                                                                          NON TER 1 1 SEQUENCE 17 AA; 1837 MW; E35DE1561000FFDC CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 AA; 2346 MW; SA282DC334CEB5EF CRC64;
                                                           01-MAY-2000 (TrEMBLrel. 13, Last Sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Putative proline-rich protein (Fragment).
Name-ctd5;
17 AA
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PRT;
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                                         01-MAY-2000 (TrEMBLrel, 13,
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Matches 4; Conservative
  PRELIMINARY;
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10 RLFLTFG 16
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Matches 4; Conservative
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8 AFVTL 12
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Q26825
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MEDLINE 87283915; PubMed=3612801;
Timmers H.Th.M., De Lange T., Kooter J.M., Borst P.;
"Coincident multiple activations of the same surface antigen gene in Trypanosoma brucei.";
Trypanosoma brucei.";
J. Mol. Biol. 194:81-90(1987).
EMBL; X05267; CAA28883.1; --
EMBL; X05267; CAA28883.1; --
EMBL; X05267; CAA28882.1; --
EMBL; X05267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bukaryota; Buglanozoa; Kinetoplastida; Trypanosomatidae; Trypanosoma
NCBI_TaxID=5691;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AF17 D (Fragment).
Ascaris suum (big roundworm) (Ascaris lumbricoides).
Eukaryota: Metazoa; Nematoda; Chromadorea; Ascaridida; Ascaridoidea;
Ascarididae; Ascaris.
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GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0019028; C:viral capaid; IEA.
GO; GO:0019031; C:viral envelope; IEA.
GO; GO:0005198; F:structural molecule activity; IEA.
INTERPRO IPR011056; Pept_824_826_C.
InterPro; IPR011056; Pept_824_826_C.
InterPro; IPR011056; Pept_824_826_C.
AIDS; Coat protein; Envelope protein; Glycoprotein; Transmembrane.
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Pred. No. 2e+03;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                                       Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nanda J.C., Stretton A.;
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases
EMBL; AY386839; AAQ90112.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE 13 AA; 1531 MW; 18DA23119D6C79C4 CRC64;
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25 AA; 2801 MW; 25E1B150CD7C14B6 CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
VSG (14 AA) (Fragment).
Trypanosoma_brucei.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
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Best Local Similarity 50.0%;
Matches 4; Conservative
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2 RNFMNFGK 9
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Q27373
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OG 06TU17
OD TO 06-TU1
DT 05-TU
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Bernards A., van der Ploeg L.H.T., Gibson W.C., Leegwater P.,
Bijgenraam F., De Lange T., Weijers P., Calafat J., Borst P.;
Rapid change of the repertoire of variant surface glycoprotein genes in trypanosomas by gene duplication and deletion.";
J. Mol. Biol. 1901-10(1986).
EMBL; X04041; CAA27674.1; -.
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NCBI_TaxID=5691;
                                                        Gaps
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PEDLINE-31139577; PubMed=1995588;

Tao T., Bourne J.C., Blumenthal R.M.;

"A family of regulatory genes associated with type II restriction-
modification systems.";

J. Bacteriol. 173:1367-1375 (1991).
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Enterobacteriaceae; Escherichia.
                                                                                                                                                                                                                                                                                                                                                                             01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 19, Last annotation update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Variant surface glycoprotein C-terminus (1 is 2nd base in codon)
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53.8%; Score 21; DB 2; Length 14;
80.0%; Pred. No. 2.2e+03;
ive 1; Mismatches 0; Indels
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SEQUENCE 15 AA; 1658 MW; DASAF6569E9A5788 CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
C protein (Fragment).
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                                                                                                                                                                                                                                                                                                                          PRELIMINARY;
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Q6EML4

Q6EML4

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Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots, rosids,
eurosids I; Cucurbitales, Cucurbitaceae, Momordica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Citrullus lanatus (Watermelon) (Citrullus vulgaris).
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons; core eudicots; rosids,
eurosids I; Cucurbitales; Cucurbitaceae; Citrullus.
                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=22951019; PubMed=14526110; DOI=10.1104/pp.103.027714; MEDLINE=2295109; PubMed=14526110; DOI=10.1104/pp.103.027714; Ayre B.G., Blair J.E., Turgeon R.; Punctional and phylogenetic analyses of a conserved regulatory program in the phloem of minor veins."; Plant Physiol. 133:1229-1239(2003). EMBL; AN379780; AQ74882.1; -.. GO; GO:0047216; F:inositol 3-alpha-galactosyltransferase acti... GO; GO:0016757; F:transferase activity, transferring glycosyl... SEQÜENCE 23 AA; 2444 MW; 62411699CAB81657 CRC64;
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Pred. No. 3.6e+03;
1; Mismatches 0; Indels
                           05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Galactinol synthase (EC 2.4.1.123) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Galactinol synthase (EC 2.4.1.123) (Fragment).
                                                                                                                                                                             Momordica charantia (Bitter gourd) (Balsam pear)
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80.0%;
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Matches 4; Conservative
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Matches 4; Conservative
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18 RAYVT 22
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RAYVT 23
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25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Aldolase B (Fragment).
Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.

PubMed=15140948; DOI=10.1093/molbev/msh157;

Axelson E., Smith N.G., Sundstrom H., Berlin S., Ellegren H.;

Male-biased mutation rate and divergence in autosomal, z-linked and w-linked introns of chicken and Turkey.";

Mol. Biol. Evol. 21:1538-1547(2004).

EMBL. AY139847; AAN75284.1;
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PubMed=15140948; DOI=10.1093/molbev/msh157;

Axelsson E., Smith N.G., Sundstrom H., Berlin S., Ellegren H.;

"Male-biased mutation rate and divergence in autosomal, z-linked and w-linked introns of chicken and Turkey.";

Mol. Biol. Evol. 21:1538-1547(2004).

EMBL, AX139841; AAN75280-1; -.
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25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Aldolase B (Fragment)
Meleagris gallopavo (Common turkey).
Meleagris gallopavo (Common turkey).
Archosauria; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Arsi Neognathae; Galliformes; Phasianidae; Meleagris.
Nisl_TaxID=9103;
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Pred. No. 2.6e+03;
1; Mismatches 1; Indels
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Pred. No. 2.6e+03;
1; Mismatches 1; Indels
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SEQUENCE 17 AA; 1813 MW; E6CF8FF0BAFA8858 CRC64;
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17 AA; 1813 MW; E6CF8FF0BAFA8858 CRC64;
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nes 4; Conservative
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Rhabdoviridae; Vesiculovirus.
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Vesicular stomatitis virus.
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                                      NCBI_TaxID=11276;
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Podatarive stress response and characterization of the oxyR-ahpC and CharakterG loci in Mycobacterium marinum.";
J. Bacteriol. 180:4856-4864(1998).
EMBL; U43810; AAC61663.1; -.
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                                                                                                      Trypanosoma cruzi.
Eukaryota, Euglenozoa, Kinetoplastida, Trypanosomatidae, Trypanosoma
                                                                                                                                                                          MEDITNE=9825151; PubMed=9556557; DOI=10.1074/jbc.273.18.10843; Di Noia J.M., D'Orso I., Aslund L., Sanchez D.O., Frasch A.C.; Di Noia J.M., D'Orso I., Aslund E., Sanchez D.O., Frasch A.C.; genes trypanosoma cruzi mucin family is transcribed from hundreds of genes having hypervariable regions."; J. Biol. Chem. 273:10843-10850(1998).

EMBL. AF036447; ARC14246.1; -.
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Bacteria, Actinobacteria, Actinobacteridae; Actinomycetales,
Corynebacterineae; Mycobacteriaceae; Mycobacterium.
NCBI_TaxID=1789;
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Pred. No. 3e+03;
1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                             51.3%; Score 20; DB 2; Length 9; 50.0%; Pred. No. 1:6e+06; tive 2; Mismatches 1; Indels
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
                               01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
                                                                                                                                                                                                                                                                     9 AA; 896 MW; DBA831B1BB5DD72D CRC64;
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        AA 6
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        PRT;
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MEDLINE=98406038; PubMed=9733688;
                                                                         Mucin-like protein (Fragment).
Name=EMUCe-19c8;
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Best Local Similarity 66.7%;
Matches 4; Conservative 1
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Matches 3; Conservative
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        PRELIMINARY;
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ALLTIG 7
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                                                                                                                                                                                                                                                                                                                                                                              4 AYTTLG 9
                                                                                                                               NCBI_TaxID=5693;
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Vesivirus.
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         SEQUENCE FROM N.A.
MEDLINE=80001959; PubMed=89911; DOI=10.1016/0092-8674(79)90274-5;
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Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AR357011; AAK43704.1; -.
NON_TER 19 19
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51.3%; Score 20; DB 2; Length 19;
Best Local Similarity 100.0%; Pred. No. 4.9e+03;
Matches 4; Conservative 0; Mismatches 0; Indels
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O91SF3;
0-1DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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RESULT 24
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01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Agnoprotein (Fragment).
Polyomavirus BK (BKV).
Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
NCBI_TAXID=10629;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=22379369; PubMed=12490781;
Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D.,
Kopp J.B.;
Transplantation 74:1497-1504 (2002).
EMBL, AF442900; AAL78924.1; -.
EMBL, AF442900; AAL78924.1; -.
EMBL, Proposition Prolyoma agno.
Interpro: IPR002643; Polyoma agno.
Pfam; PF01736; Polyoma agno; 1.
                                                                                                                                                                                                                     "Genetic and blochemical properties of an extracellular neutral metalloprotease from Staphylococcus hyicus subsp. hyicus."; Mol. Gen. Genet. 242:421-430(1994).

-!- CATALYTIC ACTIVITY: PREFERENTIAL CLEAVAGE OF BONDS WITH HYDROPHOBIC RESIDUES IN P1'.

-!- COPACTOR: BINDS A ZINC ATOM.

-!- SUBCELLULAR LOCATION: SECRETED.

GO, GO:0016787; F:hydrolaae activity, IEA.

GO, GO:0008237; F:metallopeptidase activity; IEA.

Hydrolase; Metal-binding; Metalloprotease; Zinc; Zymogen.
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Pred. No. 5.3e+03;
2; Mismatches 2; Indels
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                     Neutral metalloprotease (EC 3.4.24.31) (Pragment)
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
                                                          Staphylococcus carnosus
Bacteria, Firmicutes, Bacillales, Staphylococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21 AA.
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MEDLINE=94166751; PubMed=8121397;
Ayora S., Goetz F.;
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9 QASVKVGK 16
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                                                                                                 NCBI_TaxID=1281;
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Q8QYS5;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 14, Last annotation update)
01-JUN-2000 (TrEMBLrel. 14, Last annotation update)
Cathepsin B-like cysteine protease (Fragment).
Leishmania mexicana.
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae; Leishmania.
NBI_TAXID=5665;
                                                                                                                                                                                                                                                                                                                                                                        Feline calicivirus.
Viruses; ssRNA positive-strand viruses, no DNA stage; Caliciviridae;
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MEDLINE=94187801; PubMed=8139620; DOI=10.1016/0166-6851(93)90116-F;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rice C.C., Kruger J.M., Vilnis A., Venta P.J., Maes R.K.; Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases. EMBL; AR357010; AR43702.1; -. NON TER 9. 19
                     Length 19;
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                                                          Indels
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80.0%; Pred. No. 5.1e+03;
ive 1; Mismatches 0; Indels
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01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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100.0%; Pred. No. 4.9e+03;
tive 0; Mismatches 0;
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Pred. No. 4.9e+03;
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            Query Match
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Matches 4; Conservative
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01-JUN-2003 (
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Q91SF7;
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RESULT 26 Q09166 ID Q0916 AC Q0916 DT 01-NO DT 01-JU

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SIRAINW=UK/4A;

Adalagan J.E., Calvo S.E., Borkovich K.A., Selker E.U., Read N.D.,

Jaffe D., Fitzhugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B.,

Jaffe D., Fitzhugh W., Mang S., Nielsen C.B., Burler J., Endrizzi M.,

Blkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M.,

A Gui D., Janakiev P., Pedersen D., Nelson M., Washburne M.,

Rothe G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D.,

Kothe G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D.,

Kamal M., Kamysselis M., Mauceli E., Bielke C., Rudd S., Frishman D.,

Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S.,

Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S.,

A Gogoni C., Macino G., Catcheside D., Li W., Peratt R.J., Osmani S.A.,

DeSouza C.C., Glass L., Orbedh M.J., Berglund J., Voelker R.,

Narden O., Plamann M., Seiler S., Dunlap J., Radford A., Aramayo R.,

Natvig D.O., Alex L.A., Mannhaupt G., Ebbole D.J., Freitag M.,

Paulsen I., Sachs M.S., Lander E.S., Nusbaum C., Birren B.,

T., The Genome Sequence of the Filamentous Fungus Neurospora crassa.",
Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
NCBI_TaxID=10629;
                                                         SEQUENCE FROM N.A.
MEDLINE=22379369; PubMed=12490781;
Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D.,
Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D.,
Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D.,
Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D.,
"BK virus and SV40 co-infection in polyomavirus nephropathy.";
"Transplantation 74:1497-1504(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -1- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DBJ whole genome shotgun (WGS) entry which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Neurospora crassa.
Eukaryota, Fungi, Ascomycota, Pezizomycotina, Sordariomycetes,
Sordariomycetidae, Sordariales, Sordariaceae, Neurospora.
NCBI_TaxID=5141;
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llarity 57.1%; Pred. No. 5.6e+03;
Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                    51.3%; Score 20; DB 2; Length 21; 50.0%; Pred. No. 5.3e+03; ive 2; Mismatches 2; Indels
                                                                                                                                                                                                                          Li R.-M., Kopp J.B.;
Submitted (NOV-2018) to the EMBL/GenBank/DDBJ databases.
EMBL; AF842894; AAL78920.1; -.
GO; GO:0003677; F:DNA binding; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL, AABXO1000490; EAA28866.1; -.
SEQUENCE 22 AA; 2584 MW; 59824A08F3774EAC CRC64;
                                                                                                                                                                                                                                                                                           01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
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Best Local Similarity
Matches 4; Conserv
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Name=NCU09457.1;
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Polyomavirus BK (BKV).
Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
                                           Agnoprotein (Fragment).
Polyomavirus BK (BKV).
Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
                                                                                                                      MEDLINE=223/936. PubMed=12490781;
Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D., Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D., Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D., Transplantation 74:1497-1504(2002).

EMBL, AF442896; AAL78922.1; -
EMBL, AF42896; AAL78922.1; -
EMBL, PO03677; F:DNA binding; IEA.

Interpro; IRR002643.

ProDom; P0004470; Polyoma_agno; 1.

ProDom; P0004470; Polyoma_agno; 1.
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Pred. No. 5.38+03;
2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                              Score 20; DB 2; Length 21;
Pred. No. 5.3e+03;
2; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                 21 AA; 2375 MW; E5B8EBD9AE20D4E3 CRC64;
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Last annotation update)
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Last sequence update)
Last annotation update)
                           01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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080YSB,

080YSB,

01-JUN-2002 (TrEMBLrel. 21, La

01-JUN-2003 (TrEMBLrel. 25, Le
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01-JUN-2002 (TrEMBLrel. 21,
01-OCT-2003 (TrEMBLrel. 25,
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9 QASVKVGK 16
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Best Local Similarity
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ses 4; Conserva
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                                                                                                      MCBI_TaxID=10629;
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SEQUENCE
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RESULT 29

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Length 22;

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Nicotiana plumbaginifolia (Leadwort-leaved tobacco).
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magnoliophyta; eudicotyledons, core eudicots, asterids, lamids, Solanales, Solanaceae, Nicotiana.
                                       Rattus sp. .
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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"Alterations in the phenotype of plant cells studied by NH2-terminal
amino acid-sequence analysis of proteins electroblotted from two-
dimensional gel-separated total extracts ";
Proc. Natl. Acad. Sci. U.S.A. 84:4806-4810(1987).
                                                                                                                                                                                                                                                                               MEDLINE=93379380; PubMed=8099311; DOI=10.1016/0014-5793(93)81533-6; Raditsch M., Ruppersberg J.P., Kuner T., Gunther W., Schoepfer R., Seeburg P.H., Jahn W., Witzemann V.; "Subunit-specific Dlock of cloned NMDA receptors by argiotoxin636."; SEBS Lett. 324:63-66(1993). SEQUENCE 24 AA; 2749 MW; 203D9CE281A012C7 CRC64;
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Mammalia, Butheria, Primates; Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 100.0%; Pred. No. 6e+03; 4; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ol-WAR-2004 (TrEMBLrel. 26, Created)
01-WAR-2004 (TrEMBLrel. 26, Last sequence update)
01-WAR-2004 (TrEMBLrel. 26, Last annotation update)
Protein P7 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-2001 (TYEMBLrel. 17, Created)
01-JUN-2001 (TYEMBLrel. 17, Last sequence update)
01-JUN-2001 (TYEMBLrel. 17, Last annotation update)
DJ343K2.3 (Novel protein) (Fragment)
Name=dJ621N11.4;
Homo sapiens (Human)
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NMDA receptor subunit NR2C (Fragment).
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Matches 4, Conservative
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Best Local Similarity
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                                                                                                                                                                      NCBI_TaxID=10118;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBI_TaxID=10118;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Cicereae; Cicer.
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Pred. No. 6e+03;
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Rajeah P.N., Gupta V.S., Ranjekar P.K., Muehlbauer F.J.;
Submitrted (SEP-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AF4164811, AAL08018.1; -C.
GO, GO:0016740; F:transferase activity; IEA.
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01-UDN-2003 (TrEMBLrel. 24, Last annotation update)
Hydroxymethyl transferase (Fragment)
Cicer arietinum (Chickpea) (Garbanzo).
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(TrEMBLrel. 21, I
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01-DEC-2001 (TrEMBLrel. 19,
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8 KQFPTIG 14
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01-MAY-2000
01-MAY-2000
01-JUN-2002
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SEQUENCE
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Q9QW22;
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0945F1
10 0945F
AC 0945F
DT 01-DE
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OC EUKAR
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Q9QW23
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SEQUENCE FROM N.A.
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NCBI_TaxID=206062;
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       NCBI_TaxID=9606;
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Nature O.O.O(2003).

C.I. CAUTION: The sequence shown here is derived from an Embl. General Action of Embl. General Action (WGS) entry which is Embl. Action of States Action (WGS) entry which is
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01-JUN-2000 (TrEWBLrel. 14, Last annotation update)
Tartrate-resistant acid phosphatase PEAK 2 isoform 23 kDa subunit
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Neurospora crassa.
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Sordariomycetidae; Sordariales; Sordariaceae; Neurospora.
NCBI_TaxID=5141;
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Pred. No. 6.3e+03;
1; Mismatches 3; Indels
                                                                                                                                   Length 12;
                                                                                                                                                                                  Indels
Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AL031659; CAC34516.1; -.
NOW TER 12 12
SEQÜENCE 12 AA; 1422 MW; DC7FBF1578B2C9D2 CPC64.
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EMBL; AABX01000214; EAA32994.1; -.
SEQUENCE 14 AA; 1673 MW; 6865AEB1F564FBD4 CRC64;
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Last sequence update)
Last annotation update)
                                                                                                                                   Score 19; DB 2; L. Pred. No. 5.4e+03;
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01-MAY-2000 (TrEMBLrel. 13, Last seq
01-JUN-2000 (TrEMBLrel. 14, Last ann
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50.0%;
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                                                                                                                                        48.78;
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Best Local Similarity 50.0
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Predicted protein.
Name=NCU06392.1;
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                                                                                                                                                                                                                                                                 FVTI 9
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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DOI=10.1128/AEM.69.9.5104-5114.2003;
Rakonjac J., Ward L.J., Schiemann A.H., Gardner P.P., Lubbers M.W.,
Janckila A.J., Latham M.D., Lam K.W., Chow K.C., Li C.Y., Yam L.T., "Heterogeneity of hairy cell tartrate-resistant acid phosphatase."; Clin. Blochem. 25:437-443(1992).
SEQUENCE 16 AA; 1616 MW; C260E1A756E83299 CRC64:
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Viruses; dsDNA viruses, no RNA stage; Caudovirales; Siphoviridae;
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                                                                                                                                                              48.7%; Score 19; DB 2; Length 16; 60.0%; Pred. No. 7.1e+03; ive 1; Mismatches 1; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               replication in lactococcal prolate phages.";
Appl. Environ. Microbiol. 69:5104-5114(2003).
Bmbl; AY129506; AANUS7708.1; ---
SEQUENCE 18 AA, 2061 NW; 40DE6E393961EC06 CRC64;
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Last annotation update)
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01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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Conservative 0; Mismatches
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01-MAY-2000 (TrEMBLrel. 13,
01-MAR-2003 (TrEMBLrel. 23,
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                                Best Local Similarity
Matches 4; Conserv
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Matches
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NCBI_TaxID=3702,
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Polyomavirus BK (BKV).
Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
Vil_TaxID=10629;
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MEDLINE-22379369; PubMed=12490781;
LIR R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D.,
KOPP J.B.;
BK virus and SV40 co-infection in polyomavirus nephropathy.";
Transplantation 74:1497-1504 (2002).
Transplantation 74:1497-1504 (2002).
GBLJ, AR42903, AAL78925.1;
GO; GO:0003677; F:DNA binding; IEA.
InterPro; IPR002643; Polyoma_agno.
Pfam; PP01736; Polyoma_agno.
ProDom; P0004470; Polyoma_agno; 1.
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                                                                                                                                                        Score 19; DB 2; Length 20;
Pred. No. 8.6e+03;
1; Mismatches 1; Indels
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Graham S.W., Reeves P.A., Burns A., Olmstead R.G.;
Graham S.W., Reeves P.A., Eurns A., Olmstead R.G.;
Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AY007458, AAG12346.1;
GO; GO:0009507; C:chloroplast; IEA.
                                                                                258CC8CAA95F12D6 CRC64;
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Last sequence update)
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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Arabidopsis thaliana (Mouse-ear cress).
Chloroplast.
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            1 1
20 20
20 AA; 2223 MW;
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Best Local Similarity 60.0%;
Matches 3; Conservative
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21 AA; 2195 MW;
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Matches 3; Conservative
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8 RGFIT 12
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TVGK 21
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Q8HS54;
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08HS54
AC 08HS5
DT 01-MA
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080YS3
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Agnoprotein (Fragment).
Polyomavirus BK (BKV).
Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
NCBI_TaxID=10629;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2003 (TrEMBLrel. 25, Last annotation update)
Agnoprotein (Fragment).
Polyomavirus BK (BKV).
Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
NCBI_TaxID=10629;
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MEDILINE=22379369; PubMed=12490781;
MEDILINE=22379369; RubMed=12490781;
MEDILINE=22379369; Rirk A.D.,
Kopp J.B.,
Wannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D.,
Kopp J.B.,
We virus and SV40 co-infection in polyomavirus nephropathy.";
Transplantation 74:1497-1504(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
MEDLINE=22379369; PubMed=12490781;
MEDLINE=22379369; Robert D., Teokos M., Bynum M., Kirk A.D., La R.M., Mannon R.B., Kleiner D., Teokos M., Bynum M., Kirk A.D., Kopp J.B.;
"BK virus and SV40 co-infection in polyomavirus nephropathy.";
"Transplantation 74:1497-1504(2002).
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   DB 2; Length 21; 9e+03;
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Pred. No. 9e+03;
2; Mismatches 2; Indels
                                                             2; Indels
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Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
BMBL, A8442895, AAL798921.1;
GO; GO:0003677; F:DNA binding; IEA.
InterPro; IPR002643; Polyoma_agno.
Propom; PD00477; Polyoma_agno; 1.
Propom; PD00470; Polyoma_agno; 1.
SNOW TER 21 21
SNOW TER 21 21
SNOW TER 21 AA; 2361 MW; 65D57BD9AB20D4F6 CRC64;
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Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AR442893; AAL78919.1; -
GO, GO:0003677; F:DNA binding; IEA.
InterPro; IPR002643; Polyoma_agno.
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19; Score 19; DB ilarity 50.0%; Pred. No. 9e+C Conservative 2; Mismatches
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Polyomavirus BK (BKV).
Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
NCBI_TaxID=10629;
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MEDLINE-22379369; PubMed=12490781;
MEDLINE-22379369; Risher D., Tsokos M., Bynum M., Kirk A.D.,
Li R.M., Mannon R.B., Kleiner D., Tsokos M., Bynum M., Kirk A.D.,
Kopp J.B.;
We Virus and SV40 co-infection in polyomavirus nephropathy.";
Transplantation 74:1497-1504(2002).
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                                                                                                          Query Match

48.7%; Score 19; DB 2; Length 21;

Best Local Similarity 50.0%; Pred. No. 9e+03;

Matches 4; Conservative 2; Mismatches 2; Indels
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Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, FA442892, AAL78918.1;
EMBL, FA442892, AAL78918.1;
EMBL, FA442892, AAL78918.1;
InterPro, 1PR002643; Polyoma_agno.
Pfam; PF01736; Polyoma_agno; 1.
ProDom; P004470; Polyoma_agno; 1.
NON_TER 21
SEQÜENCE 21 AA, 2361 MW; 65D57BD9AB20D4F6 CRC64;
Pfam; PF01736; Polyoma_agno; 1.
Probom; PD004470; Polyoma_agno; 1.
21 2.
SEQÜENCE 21 AA; 2361 MW; 65D57BD9AE20D4F6 CRC64;
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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Best Local Similarity 50.0
Matches 4; Conservative
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QBQYT0
ID QBQYT0
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Search completed: May 16, 2005, 08:10:01 Job time : 170 secs

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1 RAFVTIGK 8

8 8

GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on:

May 16, 2005, 07:55:20 ; Search time 37 Seconds (without alignments) 20.804 Million cell updates/sec

1 rafvtigk 8 Title: Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

Searched:

4989 Total number of hits satisfying chosen parameters:

283416 segs, 96216763 residues

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 79:\*
1: pir1:\*
2: pir2:\*
3: pir3:\* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STIMMARTES

		d			SUMMARIES	
Result No.	Score	Query Match	Length	DB	ID	Description
-	36	92.3	20	7	865399	immunodeficiency v
7	22	56.4	25	~	B44524	pregnancy-specific
m	19	48.7	10	~	D28027	protein P7 - curle
4	19	48.7	12	N	F20907	윤
S	19	48.7	14	~	S11074	alcohol dehydrogen
9	19	48.7	22	7	B90996	probable transcrip
7	19	48.7	22	~	PT0052	translation initia
80	18	46.2	12	~	S11286	exo-alpha-sialidas
6	18	46.2	15	~	S21238	hydrogensulfite re
10	18	46.2	15	~	PA0106	protein QF200076 -
11	18	46.2	20	~	PL0145	carbon-monoxide de
12	18	46.2	21	7	A20359	translation elonga
13	18	•	23	~	843289	cytochrome-c oxida
14	18	46.2	24	N	S47563	nucleoside-diphosp
15	18	•	24	N	PX0038	methemoglobin redu
16	17	43.6	10	~	G58501	48K bile/gallbladd
17	17	•	12	~	S10626	lipovitellin - Afr
18	17	43.6	13	~	S63492	dissimilatory sulf
	17	43.6	14	7	PH1347	Ig heavy chain DJ
, 20	17	43.6	14	~	PA0109	porin por 1B - Ara
21	17	٠	14	~	PA0045	'
22	17	•	15	~	S13973	chlorophyll a/b-bi
23	17	43.6	17	N	A46592	lactase-phlorizin
24	17	٠	18	N	C56046	urinary tract ston
25	17	•	20	N	PL0161	hemagglutinin - In
56	17	٠	20	~	803505	T-cell receptor al
27	17	•	20	~	805411	carboxylesterase (
28	17	43.6	20	7	76	T-cell surface gly
59	17	43.6	21	~	149414	gene CTLA-1 protei

T-cell receptor J-mannose-1-phosphat hypothetical prote T-cell receptor al hemoglobin. extrac	pyruvate, phosphat pyruvate, phosphat protoporphyrinogen enterotoxin C-1 omega-gliadine 2' omega-gliadine 1 a	cytochrome P450 AL translation initia hypothetical prote NADH2 dehydrogenas gene RB1 protein - dihydrolipoamide S
S47202 A47415 T46622 S07729	S61410 S65629 A60757 PN0151	B61597 JN0264 D83865 T14224 I78870 A37823
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2222 18447	221111 212244	14 16 16 17
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43.6	24444 21444 20000	44444 00000000
11 11 11 11	16.5 16 16 16 16	9110
3 3 3 3 4 4	33333 334 337 337	044444 01224 012643

## ALIGNMENTS

RESULT 1 S65399 Cimmunodeficiency virus type 1, HIV-1 gpl20 - human (fragments) CiSpecles: Homo sapiens (man) C;Specles: Boot-1996 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999 C;Accession: S65399 R;Niwa, Y:, Yano, M.; Futaki, S.; Okumura, Y.; Kido, H.
 A;Title: T-cell membrane-associated serine protease, tryptase TL(2), binds human immunod man immunod faiteernce number: S65399; MUID:96203909; PMID:8620895 A;Reference number: S65399 A;Retesion: S65399 A;Reternce preliminary A;Molecule type: protein A;Residues: 1-10;11-20 cMIW> C;Superfamily: type B retrovirus env polyprotein
 Query Match 92.3%; Score 36; DB 2; Length 20; Best Local Similarity 87.5%; Pred. No. 0.11; Batches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

|||||||| RAFVTIGR 12 1 RAFVTIGK 8

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pregnancy-specific glycoprotein SBU-3-62 - sheep (fragment)
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Species: 30-Sep-1993 #sequence\_revision 30-Sep-1993 #text\_change 16-Aug-2004
C;Accession: 184524
A;Atkinson, Y.H.
submitted to the Protein Sequence Database, June 1993
A;Reference number: A44524
A;Accession: B44524
A;Status: preliminary
A;Molecule type: protein
A;References: UNIPROT:Q9TRE1
C;Superfamily: Pepsin
C;Keywords: glycoprotein Length 25; Query Match 56.4%; Score 22; DB 2; Length 25; Best Local Similarity 57.1%; Pred. No. 2.6e+02; Matches 4; Conservative 1; Mismatches 2; Indels B44524

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Gaps

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A;Cross-references: UNIPROT:Q8X365; GB:BA000007; PIDN:BAB36361.1; PID:g13362407; GSPDB:GP:A;Experimental source: strain 0157:H7, substrain RIMD 0509952
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           riving translation initiation factor eIF-2 gamma chain - pig (fragment)
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 09-Jul-2004
C;Accession: PT0052
A;Risuki, H.; Mukouyama, E.B.
Agric. Biol. Chem. 52, 1397-1408, 1988
A;Ritle: Pigl liver translational initiation factor eIF-2: N-terminal amino acid sequences
A;Reference number: PT0051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic Acids Res. 6, 3745-3757, 1979
A;Title: 5' and 3' terminal nucleotide sequences of the RNA genome segments of influenza
A;Reference number: S11286; MUID:80034428; PMID:493121
A;Accession: S11286
                                                          probable transcription regulator [imported] - Escherichia coli (strain 0157:H7, substrain C;Species: Escherichia coli (C;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 09-Jul-2004
                                                                                                                                                                      C;Accession: B90996
R;Hayashi, T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.; Han, C.G., gasawara, N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H.
DNA Res. B, 11-22, 2001
A;Fille: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7 and genome A;Reference number: A99629; MUID:21156231; PMID:11258796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   - influenza A virus (strain A/PPV/Rostock/34 [H7N1]) (
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C;Species: influenza A virus
C;Date: 31-Dec:1990 #sequence_revision 31-Dec-1990 #text_change 22-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: genomic RNA
A;Residues: 1-12 <ROB>
A;Cross-references: EMBL:J02114; NID:g324483; PIDN:AAA43398.1; PID:g324486
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Pred. No. 1.2e+03;
1; Mismatches 2; Indels
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Pred. No. 1.2e+03;
2; Mismatches 1;
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A,Residues: 1-22 <SUZ>
A,Cross-references: UNIPROT:P20461
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Similarity 57.1%;
4; Conservative 1
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Matches 4; Conservative
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AGVTLGQ 11
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Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                      A; Accession: B90996
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-22 <HAY>
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R; Robertson, J.S.
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A;Gene: ECs2938
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C;Species: Gadus morhua callarias (Baltic cod)
C;Species: Gadus morhua callarias (Baltic cod)
C;Species: 30-Jun-1991 #sequence_revision 30-Jun-1991 #text_change 31-Jan-1997
C;Accession: S11074 #.; Batchius, M.; Danielsson, O.; Persson, B.; Cederlund, E.; Kaiser, R.; Hol FEBS Lett. 269, 194-196, 1990
FFBS Lett. 269, 194-196, 1990
FFBS Lett. 269, 194-196, 1990
A;Title: East atom bombardment mass spectrometry and chemical analysis in determinations A;Reference number: S11074
A;Molecule type: Drotein
A;Molecule type: Drotein
A;Residues: 1-5;6-14 <EGE>
C;Keywords: acetylated amino end; alcohol metabolism; NAD; oxidoreductase
F;1/Modified site: acetylated amino end (Ala) #status experimental
protein P7 - curled-leaved tobacco (fragment)
C;Species: Nicotiana plumbaginifolia (curled-leaved tobacco)
C;Species: Nicotiana plumbaginifolia (curled-leaved tobacco)
C;Species: 19-May-1989 #sequence_revision 19-May-1989 #text_change 09-Jul-2004
C;Accession: D28027
R;Bauw, G; De Loose, M.; Inze, D.; Van Montagu, M.; Vandekerckhove, J.
R;Bauw, G; De Loose, M.; Inze, D.; Van Montagu, M.; Vandekerckhove, J.
R;Title: Alterations in the phenotype of plant cells studied by NH2-terminal amino acid-A;Reference number: A94167
A;Reference number: A94167
A;Residues: 1-10 cBAU>
A;Residues: 1-10 cBAU>
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8est Local Similarity 60.0%; Pred. No. 6.3e+02;
Matches 3; Conservative 1; Mismatches 1; Indels
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66.7%; Pred. No. 5.3e+02;
tive 1; Mismatches 1;
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Best Local Similarity 66.7
Matches 4; Conservative
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46.2%; Score 18; DB 2; I ilarity 75.0%; Pred. No. 2.1e+03; Conservative 1; Mismatches 0;
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Best Local Similarity
Matches 3; Conserv
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Matches 3; Conserv
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A; Residues: 1-23 <MAS>
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16 ITIG 19
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$21238
hydrogensulfite reductase (EC 1.8.99.3) beta chain - Desulfovibrio vulgaris (fragment)
hydrogensulfite reductase (EC 1.8.99.3) beta chain - Desulfovibrio vulgaris (fragment)
h/Alternate names: bisulfite reductase; desulforuscidin; desulforubidin; desulfoviridin;
C;Species: Desulfovibrio vulgaris
C;Date: 19-Mar-1997 #sequence_revision 11-Jun-1999 #text_change 11-Jun-1999
C;Accession: $21238
R;Pierik, A.J.; Duyvis, M.G.; van Helvoort, J.M.L.M.; Wolbert, R.B.G.; Hagen, W.R.
A;Fitle: The third subunit of desulfoviridin-type dissimilatory sulfite reductases.
A;Reference number: $21197; MUID:92209491; PMID:1555572
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carbon-monoxide dehydrogenase (EC 1.2.99.2) small chain - Pseudomonas carboxydoflava (fr
C;Species: Pseudomonas carboxydoflava
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                                                                                          46.2%; Score 18; DB 2; Length 12; 75.0%; Pred. No. 1.1e+03; tive 1; Mismatches 0; Indels
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Pred. No. 1.4e+03;
2; Mismatches 1;
A;Map position: segment 6
C;Superfamily: influenza virus exo-alpha-sialidase
C;Keywords: glycosidase; hydrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: protein
A;Residues: 1-15 <PIE>
A;Experimental source: strain Hildenborough
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                                                                                     Query Match
Best Local Similarity 75.0
Matches 3; Conservative
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Matches 3; Conservative
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A, Molecule type: protein
A, Residues: 1-15 < CHO>
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AILTLG 9
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1 AFISSG 6
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ITIG 11
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Best Local &
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structural genes in carboxydotrop?
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A;Residues: 1-20 <KGA>
C;Comment: Carbon-monoxide dehydrogenase consists of three polypeptide chains: large, mec
C;Keywords: oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C,Accession: A20359
R,Slobin, L.I.; Clark, R.V.; Olson, M.O.J.
Biochemistry 22, 1911-1917, 1981
A;Title: Limited cleavage of eucaryotic elongation factor Tu by trypsin: alignment of th A;Reference number: A20359; MUID:83204805; PMID:6682677
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C;Date: 19-Mar-1997 #sequence_revision 01-May-1998 #text_change 09-Jul-2004
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A)Genetic code: SGC6
C;Superfamily: cytochrome-c oxidase chain III
C;Keywords: electron transfer; membrane-associated complex; mitochondrial inner membrane
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C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 28-Apr-1993 C;Accession: PL0145
R;Kraut, M.; Hugendieck, I.; Herwig, S.; Meyer, O. Arch. Microbiol. 152, 335-341, 1989
A;Title: Homology and distribution of CO dehydrogenase structural genes in C A;Reference number: PL0138; MUID:90055678; PMID:2818128
A;Accession: PL0145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                translation elongation factor EP-Tu, mitochondrial - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 09-Jul-2004
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R;Maslov, D.A.; Avila, H.A.; Lake, J.A.; Simpson, L. Nature 368, 345-348, 1994
A;Title: Evolution of RNA editing in kinetoplastid proto A;Reference number: S43286; MUID:94173338; PMID:8127370
                                                                                                                                                                                                                                                                                                                                                                                                       h 46.2%; Score 18; DB 2; I Similarity 60.0%; Pred. No. 1.8e+03; 3; Conservative 1; Mismatches 1;
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Pred. No. 1.9e+03;
1; Mismatches 0;
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A,Note: residue 19 has also been sequenced as Lys
C,Keywords: mitochondrion; protein biosynthesis
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Tipovitellin - African clawed frog
CiSpecies: Kanopus laevis (African clawed frog)
CiSpecies: Renopus laevis (African clawed frog)
CiAccession: S10626
R;Wallace, R.A.; Hoch, K.L.; Carnevali, O.
J. Mol. Biol. 213, 407-409, 1990
A;Title: Placement of small lipovitellin subunits within the vitellogenin precursor in XA; Reference number: S10624; MUD:90278951; PMID:2352275
A;Molecule type: protein
A;Residues: 1-12 < WAL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C, Accession: PH1347
R; Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A, Title: Predominance of fetal type DJH joining in young children with B precursor lymphc A; Reference number: PH1302; MUID:93094761; PMID:1460419
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
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                                                                                                  Length 10;
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Pred. No. 1.9e+03;
0; Mismatches 3; Indels
                  A, Experimental source: human bile and gallbladder stones A,Note: 1-Ser and 4-Glu were also found
                                                                                               43.6%; Score 17; DB 2; Le 66.7%; Pred. No. 1.6e+03; live 0; Mismatches 2;
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Pred. No. 2e+03;
1; Mismatches
A; Cross-references: UNIPROT: Q7M1C8
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Best Local Similarity 62.5
Matches 5; Conservative
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A Molecule type: protein
A; Residues: 1-13 <STE>
                                                                                                            Best_Local_Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 3; Conserv
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1 AFIPTG 6
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A; Residues: 1-14 < WAS>
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PX0038
methemoglobin reductase (NADPH) (EC 1.6.2.-) - bullfrog (fragment)
C;Species: Rana catesbeiana (bullfrog)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 09-Jul-2004
C;Accession: PX0038
R;Abe, Y: Ito, T: Okazaki, T.
J. Biochem. 108, 255-260, 1990
A;Title: Purification and characterization of NADPH-dependent methemoglobin reductase franchemoce number: PX0038; MUID:91035356; PMID:2172227
                                                                                                                                                                                              nucleoside-diphosphate kinase (EC 2.7.4.6) - oat (fragment)
C;Species: Avena sativa (oat)
C;Date: 07-May-1995 #sequence_revision 24-Oct-1997 #text_change 09-Jul-2004
C;Accession: 847663
R;Sommer, D.; Song, P.S.
B;Commer, D.; Song, P.S.
A;Title: A plant nucleoside diphosphate kinase homologous to the human Nm23 gene product
A;Réference number: 847563; MUID:9431244; PMID:8038216
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G58501
48K bile/gallbladder stone protein - unidentified bacterium (fragment)
48K bile/gallbladder stone protein - unidentified bacterium (fragment)
C;Species: unidentified bacterium
C;Species: unidentified bacterium
C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_change 09-Jul-2004
C;Accession: G58501
R;Blantette, M.B.
R;Dainette, J.P.; Binette, M.B.
R;Dainette to the Protein Sequence Database, October 1996
A;Description: The proteins of kidney and gallbladder stones.
A;Reference number: A58501
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C,Superfamily: nucleoside diphosphate kinase
C,Keywords: phosphotransferase; pyrimidine nucleotide biosynthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46.2%; Score 18; DB 2; Length 24; 50.0%; Pred. No. 2.2e+03; cive 1; Mismatches 2; Indels
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A;Experimental source: nucleated erythrocyte
C;Keywords: NADP; oxidoreductase
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Best Local Similarity 50.0
Matches 3; Conservative
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nes 3, Conservative
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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-24 <SOM>
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A;Residues: 1-24 <ABE>
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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-10 <BIN>
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7 ITIG 10
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17 VTLGQ 21
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                       4 VTIG 7
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C;Accession: A46592
R;Dudley, M.A.; Hachey, D.L.; Quaroni, A.; Hutchens, T.W.; Nichols, B.L.; Rosenberger, J. S. Biol. Chem. 268, 13609-13616, 1993
A;Title: In vivo sucrase-isomaltase and lactase-phlorizin hydrolase turnover in the fed A;Reference number: A46592; MUID:93293888; PMID:8514793
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R;Sweetser, M.T.; Braciale, V.L.; Braciale, T.J.
J. Exp. Med. 170, 1357-1368, 1989
A;Title: Class I major histocompatibility complex-restricted T lymphocyte recognition of
                                                   C; Keywords: chlorophyll; chloroplast; light-harvesting complex; thylakoid; transmembrane
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                                                                                                                                                                                                                                                                                                                                                                      lactase-phlorizin hydrolase, 200K isoform - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 16-Feb-1994 #sequence_revision 18-Nov-1994 #text_change 01-Nov-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    urinary tract stone matrix protein 5, 32K - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 12-Apr-1995 #sequence_revision 12-Apr-1995 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hemagglutinin - Influenza H2N2 (fragment)
C;Species: influenza H2N2
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Accession: C56046
R;Binette, J.P.; Binette, M.B.; Gawinowicz, M.A.; Kendrick, N.
submitted to the Protein Sequence Database, February 1995
A;Description: Isolation, characterization and sequence of stone proteins.
A;Reference number: A56046
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                                                                                              43.6%; Score 17; DB 2; Length 15; 60.0%; Pred. No. 2.3e+03; tive 1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-17 kDUD:
A;Note: sequence extracted from NCBI backbone (NCBIP:134559)
C;Keywords: carbohydrate digestion; intestine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               43.6%; Score 17; DB 2; I 42.9%; Pred. No. 2.6e+03; tive 1; Mismatches 3;
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A;Molecule type: protein
A;Residues: 1-18 <BIN:
A;Cross-references: UNIPROT:Q7M4Q7
                                                                                                 Query Match
Best Local Similarity 60.0°
Matches 3; Conservative
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9 RTYAAVG 15
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6 FTSIG 10
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                              A; Genome: nuclear
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     C; Genetics:
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 30-Jun-1992 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
C;Accession: PA0045
R;Kamo, M.; Kawakami, T.; Miyatake, N.; Tsugita, A.
Submitted to JIPID, July 1994
A;Description: Separation and characterization of Arabidopsis proteins by two-dimensiona
A;Reference number: PA0001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chlorophyll a/b-binding protein type II - garden pea (fragment)

cispecies: Pieum sativum (garden pea)

cispecies: Pieum sativum (garden pea)

cipactes: 19-Mar-1997 #sequence_revision 24-Mar-1999 #text_change 24-Mar-1999

cipacession: 513973

RiJahns, P.; Junge, W.

Bur. J. Blochem: 193, 731-736, 1990

A;Title: Dicyclohexylcarbodilmide-binding proteins related to the short circuit of the A;Reference number: 513973; MUID:91065379; PMID:2174365
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Accession: PAp.1995 #sequence_revision 26-May-1995 #text_change 09-Jul-2004
C;Accession: PA0109
R;Kamo, M.; Kawakami, T.; Tsugita, A.
submitted to JIPID, March 1995
A;Reference number: PA0109
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                                                43.6%; Score 17; DB 2; Length 14; 60.0%; Pred. No. 2.2e+03; tive 1; Mismatches 1; Indels
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Pred. No. 2.2e+03;
1; Mismatches 2; Indels
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A;Residues: 1-14 «KAM»
A;Crosa-references: UNIPROT:Q8LA84; UNIPROT:Q42292
A;Experimental source: root
C; Keywords: heterotetramer; immunoglobulin
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A;Mesidues: 1-14 «KAM»
A;Cross-references: UNIPROT:Q7MIW9
A;Experimental source: root
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Best Local Similarity 60.0
Matches 3; Conservative
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7 YTEIGK 12
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PA0045
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R;Classon, J. T. Tragaratos, J.; Kirszbaum, L.; Maddox, J.; Mackay, C.R.; Brandon, M.; M. Immunogenetics 23, 129-132, 1986
A;Title: The L3T4 antigen in mouse and the sheep equivalent are immunoglobulin-like.
A;Reference number: A47642; MUID:86166694; PMID:3082751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Accession: 149414
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-21 <RES>
A;Cross-references: UNIPROT:Q62538; EMBL:UO5708; NID:g497037; PIDN:AAB60471.1; PID:g4970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Ko, M.S.; Wang, X.; Horton, J.H.; Hagen, M.D.; Takahashi, N.; Maezaki, Y.; Nadeau, J.H. Mamm. Genome 5, 349-355, 1994
A;Title: Genetic mapping of 40 cDNA clones on the mouse genome by PCR.
A;Reference number: I48934; WUID:94319082; PMID:8043949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C.Species: Mus spretus (western wild mouse)
C.Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 09-Jul-2004
C.Accession: 149414
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C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 23-Jul-1999
                          T-cell surface glycoprotein CD4 - sheep (fragment).
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: B47642
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Pred. No. 3.1e+03;
1; Mismatches 1; Indels
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Pred. No. 3.2e+03;
1; Mismatches 2;
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R;Plaza, A.; Kono, D.H.; Theofilopoulos, A.N.
submitted to the EMBL Data Library, February 1993
A;Reference number: S40133
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                                                                                                                                                                                                                                                                                                                        A,Molecule type: protein
A,Residues: 1-20 cCLA-
A,Cross-references: UNIPROT:P05542
C,Keywords: glycoprotein
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Best Local Similarity 50.0%;
Matches 3; Conservative
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Best Local Similarity 60.0
Matches 3; Conservative
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A;Molecule type: mRNA
A;Residues: 1-21 <PLA>
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A;Status: preliminary
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N.Alternate names: serine esterase
C;Species: Sulfolobus acidocaldarius
C;Species: Sulfolobus acidocaldarius
C;Accession: 505411
R;Sobek, H.; Goerisch, H.
Biochem. J. 261, 993-998, 1989
A;Title: Further kinetic and molecular characterization of an extremely heat-stable carb A;Reference number: S05411; MUID:90026296; PMID:2508625
A;Accession: S05411
                    A;Accession: PL0161
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-20 cSWE>
A;Esperimental type: mRNA
A;Evperimental source: strain A/JAP/305/57
A;Experimental source: strain A/JAP/305/57
C;Comment: This protein plays a major role in initiation of infection and in the pathoge C;Superfamily: influenza virus hemagglutinin
C;Keywords: hemagglutinin
F;1-20/Region: immunodominant site recognized by T-lymphocytes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T-cell receptor alpha chain J region (80) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change 30-May-1997
C;Accession: S03505
R;Winoto, A.; Mjolsness, S.; Hood, L.
Nature 316, 832-836, 1985
A;Title: Genomic organization of the genes encoding mouse T-cell receptor alpha-chain.
A;Reference number: S03503; MUID:85296332; PMID:2993908
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40.0%; Pred. No. 3.1e+03;
tive 3; Mismatches 0; Indels
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A;Molecule type: DNA
A;Cross-references: EMBL:X02859
A;Note: this sequence was determined from the germline gene
C;Keywords: T-cell receptor
A; Reference number: PL0161; MUID: 90010790; PMID: 2477491
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Best Local Similarity 60.0%; Pred. No. 3.1e+03;
Matches 3; Conservative 1; Mismatches 1;
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Pred. No. 3.1e+03;
1; Mismatches 1;
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A;ONOTE: 1-Ala and 1-Ser were also found
C;Keywords: carboxylic ester hydrolase; tetramer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               43.6%;
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                                                                                                                                                                                                                                                                                                                                                             2; Conservative
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Best Local Similarity
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10 YVSVG 14
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C,Accession: S65729

R,Fushitani, K.; Higashiyama, K.; Asao, M.; Hosokawa, K.
Bjochim. Blophya. Acta 1292, 273-280, 1996

A;Title: Characterization of the constituent polypeptides of the extracellular hemoglobi. A;Reference number: S65721; MUID:96176855; PMID:8597573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pyruvate, phosphate dikinase (BC 2.7.9.1), cytosolic - Flaveria trinervia (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hemoglobin, extracellular, chain d2 - earthworm (Lumbricus terrestris) (fragment)
C;Species: Lumbricus terrestris (common earthworm)
C;Date: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Flaveria trinervia
C;Date: 27-Apr-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
                                                                                                                                            A;Cross-references: EMBL:M38675; NID:g201207; PIDN:AAA40193.1; PID:g201208
A;Experimental source: strain Balb/c
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                              mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
Nucleic Acids Res. 16, 11355-11364, 1988
A;Title: The structure of V-alpha and J-alpha segments in the
A;Reference number: S06466; MUID:89083566; PMID:2849763
                                                                                                                                                                                                                                                                                                                                                 Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 25;
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                                                                                                                                                                                                                      A,Map position: 14
C;Keywords: glycoprotein; heterodimer; T-cell receptor
F;1-22/Domain: J segment <JSE>
F;23-24/Domain: C region (fragment) <CRE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16.5; DB 2;
Pred. No. 4.2e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                 43.6%; Score 17; DB 2; 1 60.0%; Pred. No. 3.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17; DB 2; I
Pred. No. 3.9e+03;
0; Mismatches 1;
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C;Keywords: oxygen carrier
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Best Local Similarity 62.5%;
Matches 5; Conservative
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Best Local Similarity 80.0%;
Matches 4; Conservative
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Best Local Similarity
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A; Residues: 1-24 < YAG>
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                                                                                                                                                                                                                                                                                                                                                                                                                                               4 VTIGK 8
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                                                                                                                                                                                                                   mannose-1-phosphate guanylyltransferase (EC 2.7.7.13) 37K beta chain - pig (fragment)
N'Alternate names: GDP-mannose pyrophosphorylase 37K beta chain
C;Species Sus scrofa domestica (domestic pig)
C;Date: 25-Feb-1994 #sequence_revision 12-Aug-1996 #text_change 09-Jul-2004
C;Accession: A47415
R;Szumilo, T.; Drake, R.R.; York, J.L.; Elbein, A.D.
J. Biol. Chem. 268, 17943-17950, 1993
A;Title: GDP-mannose pyrophosphorylase. Purification to homogeneity, properties, and uti
A;Reference number: A47415; MUID:93352609; PMID:7688733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Accession: T46622
R;Chang, S.; Puryea, J.; Funkhouser, E.A.; Newton, R.J.; Cairney, J.
Bubmitted to the EMBL Data Library, July 1995
A;Reference number: Z23105
A;Reference number: Z23105
A;Reference number: Z23105
A;Recession: T46622
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Residues: 1-24 cCHA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Description: generates GDP-mannose and pyrophosphate from mannose-1-phosphate and GTP A; Noce: also catealyzes synthesis of GDP-glucose from glucose-1-phosphate (EC 2.7.7.34 ac C; Superfamily: mannose-1-phosphate guanylyltransferase C; Keywords: nucleotidyltransferase
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T-cell receptor alpha chain J segment (DT) - mouse (fragment)
C.ppecies: Mus musculus (house mouse)
C.pate: 29-Jan-1993 #sequence_revision 29-Jan-1993 #text_change 05-Nov-1999
C.Accession: S07699
R.Yague, J.; Blackman, M.; Born, W.; Marrack, P.; Kappler, J.; Palmer, E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C, Species: Pinus taeda (loblolly pine)
C, Date: 18-Feb-2000 #sequence_revision 18-Feb-2000 #text_change 18-Feb-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT: 09TRF4
A;Note: sequence extracted from NCBI backbone (NCBIP:136438)
C;Complex: The enzyme appears to be a heterodimer of alpha and beta chains.
C;Function:
       Gaps
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     1; Indels
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A;Experimental source: strain s6PT2x86PT3; 8 month seedlings
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     1; Mismatches
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     3; Conservative
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nes 4; Conserva
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KALILVG 8
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LTFGK 13
                                                4 VTIGK 8
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Best Local &
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cytochrome P450 AL-2 - rat (fragment)
cytochrome P450 AL-2 - rat (fragment)
c;pacies: Rattus norvegicus (Norway rat)
c;pate: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
c;Accession: 861597
R;Shimeno, H.; Toda, A.; Ogata, S.; Nagamatsu, A.
R;Shimeno, H.; Toda, A.; Ogata, S.; Nagamatsu, A.
A;Title: Purification and aminopytine enonoxygenase activity of liver microsomal cytochromater number: A61597; MUID:91292910; PMID:1676625
                                                                                                                                                                                                                                                                                                                omega-gliadine 1 and 2 - Aegilops longissima (fragment)
C:Species: Aegilops longissima
C:Species: Aegilops longissima
C;Accession: PN0147; PN0146
R;Odintsova, T.I.; Egorov, T.A.
Biokhimia 55, 509-516, 1990
A;Rttle: N-Terminal sequences of omega-gliadins of Aegilops longissima: On the origin of A;Reference number: PN0146; MUID:90283493; PMID:2354218
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N;Alternate names: eIF2 gamma chain
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 09-0ct-1992 #sequence_revision 09-0ct-1992 #text_change 09-Jul-2004
C;Accession: JN0264
R;Mukouyama, B.B.; Shiokawa, H.; Suzuki, H.
Biosci. Biotechnol. Biochem. 56, 680-681, 1992
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                                     4; Indels
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50.0%; Pred. No. 3.7e+03; ive 0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Accession: PN0146
A;Molecule type: protein
A;Residues: 1-9 <0D2>
A;Experimental source: strain K-202
A;Experimental source: strain K-202
A;Note: omega-gliadine 1 (amino-terminal fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A Cross-references: UNIPROT:Q7M1V5
A;Experimental source: strain K-202
A;Note: omega-gliadine 2 (amino-terminal fragment)
A;Accession: PN0146
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Best Local Similarity 50.0
Matches 4; Conservative
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Best Local Similarity 33.3
Matches 2; Conservative
                                     4; Conservative
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A,Molecule type: protein
A,Residues: 1-14 <SHI>
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Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 AFVTIG 7
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                                                                                                             1 RAFVTIGK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 40
B61597
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C;Species: Aegilops longissima
C;Species: Aegilops longissima
C;Species: 30-Sep-1933 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PN0151
R;Odintsova, T.I.; Egorov, T.A.
B;Odintsova, T.I.; Egorov, T.A.
B;Odintsova, T.I.; Egorov, T.A.
B;Odintsova, T.I.; Egorov, T.A.
B;Odintsova, PN0151
B;Odintsova, PN0151
B;Odintsova, T.I.; Egorov, T.A.
B;Odintsova, T.I.; Egorova, T.A.
B;Odintsova, T.A.
B;Odintsova
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C;Species: Staphylococcus aureus
C;Species: Staphylococcus aureus
C;Date: 14-May-1993 #sequence_revision 14-May-1993 #text_change 30-Sep-1993
C;Date: 14-May-1993 #sequence_revision 14-May-1993 #text_change 30-Sep-1993
C;Accession: A60757
R;Bohach, G.A.; Handley, J.P.; Schlievert, P.M.
Infect. Immun. 57, 23-28, 1989
Infect. Immun. 57, 23-28, 1989
A;Title: Biological and immunological properties of the carboxyl terminus of staphylocochy, Reference number: A60757; MulD:89079292; PMID:2909489
A;Accession: A60757
A;Recession: Decliminary
A;Residues: 1-12 < BOH>
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R;Taketani, S.; Yoshinaga, T.; Furukawa, T.; Kohno, H.; Tokunaga, R.; Nishimura, K.; Inc Bur, J. Blochem. 230, 760-765, 1995
A;Title: Induction of terminal enzymes for heme biosynthesis during differentiation of m A;Reference number: S65629; MUID:95331315; PMID:7607249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A, Pathway: heme biosynthesis, porphyrin biosynthesis
C, Superfamily: phytoene dehydrogenase
C, Keywords: heme biosynthesis; mitochondrion; oxidoreductase; porphyrin biosynthesis
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                                                                                                                                                                                                                             protoporphyrinogen oxidase (EC 1.3.3.4) - bovine (fragment)
C;Species: Bos primigenius taurus (cattle)
C;Date: 14-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 26-May-2000
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80.0%; Pred. No. 3.2e+03;
.ive 0; Mismatches 1; Indels
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Best Local Similarity 80.0
Matches 4; Conservative
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                  7 VTGGK 11
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Query Match

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SFLLVG 14
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Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans and A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Accession: D83865
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-16 <STO>
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A;Experimental source: strain C-125
C;Genetics:
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A,Status: preliminary; translated from GB/EMBL/DDBJ
A,Molecule type: DNA
A,Molecule type: DNA
A,Rosecule type: DNA

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C; Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 09-Jul-2004
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submitted to the EMBL Data Library, May 1996
A;Description: Evolution of pulmonate gastropod mitochondrial genomes: Comparisons of co
A;Title: GTP-binding sequences in the gamma subunit of pig liver initiation factor 2. A;Reference number: JN0264; MUD:92282179; PMID:1368212
A;Accession: JN0264
A;Molecule type: protein
A;Residues: 1-16 cMUX-
A;Cross-references: UNIRROT:09TRQ9
A;Cross-references: UNIRROT:09TRQ9
C;Keywords: GTP binding #status experimental
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C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 09-Jul-2004
C;Accession: D83865
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Pred. No. 4.2e+03;
2; Mismatches 2; Indels
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C;Keywords: mitochondrion; NAD; oxidoreductase
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Best Local Similarity 42.9
Matches 3; Conservative
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Matches 3; Conservative
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Best Local Similarity 33.3
Matches 2; Conservative
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QATINIG 7
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T14224
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2 AFVTIG 7

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R;Hogg, A.; Onadim, Z.; Baird, P.N.; Cowell, J.K.
Oncogene 7, 1445-1451, 1992
A;Title: Detection of heterozygous mutations in the RB1 gene in retinoblastoma patients 1
A;Reference number: I58362; MUID:92319557; PMID:1352398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R,Rahmatullah, M.; Radke, G.A.; Andrews, P.C.; Roche, T.E.
Distol. Chem. 265, 14512-14517, 1990
A,Title: Changes in the core of the mammalian-pyruvate dehydrogenase complex upon select:
A,Reference number: A37823; MUID:90354445; PMID:2167319
                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT: Q92727; GB:L41911; NID: 9794004; PIDN:AAB59483.1; PID: 9794005
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                               C;Species: Homo sapiens (man)
C;Date: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C,Species: Bos primigenius taurus (cattle)
C,Date: 30-Apr_1991 #sequence_revision 30-Apr-1991 #text_change 09-Jul-2004
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Pred. No. 4.5e+03;
2; Mismatches 0; Indels
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Pred. No. 4.5e+03;
1; Mismatches 1;
                                                                                                                                                                                                                                      A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-17 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                           A, Cross-references: GDB:118734; OMIM:180200
A, Map position: 13q14.3-13q14.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Search completed: May 16, 2005, 08:10:44 Job time : 39 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A, Cross-references: UNIPROT:Q7M2M8
C, Keywords: acyltransferase; coenzyme A
gene RB1 protein - human (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 60.0%;
Matches 3; Conservative
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Best Local Similarity 60.0
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Accession: A37823
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-17 <RAH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 VTIGK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1:||:
3 VSIGE 7
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                                                                               C; Accession: I78870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C; Accession: A37823
                                                                                                                                                                                                                                                                                                                                                                                    A; Gene: GDB: RB1
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

- protein search, using sw model OM protein May 16, 2005, 12:53:32 ; Search time 18.0769 Seconds (without alignments) 79.839 Million cell updates/sec Run on:

US-08-869-386-1 Title: Perfect score:

1 RIQRGPGRAFVTIGK 15 Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283416 segs, 96216763 residues Searched:

4989 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

PIR 79:\*
1: pir1:\*
2: pir2:\*
3: pir3:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SIMMARIES

		immunodeficiency v	psin	por 1B	porl -	protein P7 - curle	exo-alpha-sialidas	hydrogensulfite re	dihydrolipoamide S	T-cell receptor be	Ig kappa-1 chain J	T-cell receptor be	dissimilatory sulf	biliary glycoprote	hypothetical prote	MSEL-neurophysin -	bombinin-like pept	36K protein 3124 -	cytochrome-c oxida	heterocyst-inhibit	cytochrome-c oxida	rin	82K protein - bovi	dentinal fluid tra	pyruvate dehydroge	T-cell receptor be	T-cell receptor J-	peroxidase (EC 1.1	pec	alpha-amylase inhi
SUMMARIES	CI CI	565399	548654	PA0109	PA0045	D28027	S11286	821197	A37823	PT0515	C53275	PH0915	S63490	S31427	C42856	B60422	D41575	PS0453	865388	AF2093	877991	865629	831220	DIRT	A60225	PQ0070	S47192	S22221	4	S10850
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	Result No.	-	67	e	4	ហ	ų	7	80	6	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	. 28	29

cytochrome-c oxida chaperone, TCP1-re Ig heavy chain DJ T-cell receptor all myosin light chain cystathionine beta 2S albumin small c TCR delta chain V- LX-1 tumor antigen probable gag polym Ig heavy chain J-H Ile-Ser-bradykinin cytochrome-c oxida T-cell receptor betransforming probable coxida T-cell receptor betransforming probable transforming probable transforming proceeding T-cell receptor betransforming proceeding proceeding the procedulation of the proceeding the proceeding the proceeding the proceeding the procedulation of the proceeding the proceeding the proceeding the proceeding the proceeding the procedulation of the proceeding the proceeding the proceeding the proceeding the procedulation of the proceeding the procedulation of the procedulation	T cell receptor al
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2 2 2 2 2 2 2 2 2 2 3 2 3 3 3 3 3 3 3 3	20
8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	45

## ALIGNMENTS

 RESULT 1 S65399
immunodeficiency virus type 1, HIV-1 gp120 - human (fragments) C.Species: Homo sapiens (man)
C.Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
Cjaccesion: 309339 R;Niwa, Y.; Yano, M.; Futaki, S.; Okumura, Y.; Kido, H.
Eur. J. Biochem. 237, 64-70, 1996 A;Title: T-cell membrane-associated serine protease, tryptase TL(2), binds human immunodk
man immunodeficiency virus type 1 inhibit cleavage of gp120.
A;Accession: 665399
A;Status: preliminary
A;Molecule type: protein
A; Residues: 1-10;11-20 <niw></niw>
C;Superfamily: type E retrovirus env polyprotein
 Query Match 77.9%; Score 60; DB 2; Length 20;
 Best Local Similarity 91.7%; Pred. No. 0.00082; Matched 11. Canadariting 1. Missatched 0. Indels 0. Gans 0.
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 Qy 4 RGPGRAFVTIGK 15

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RESULT 2
848654
Plasmedium falciparum)
C;Species: Plasmodium falciparum)
C;Species: Plasmodium falciparum
C;Date: 15-Jul-1995 #sequence\_revision 19-Apr-1996 #text\_change 09-Jun-2000
C;Accession: S48654 #sequence\_revision 19-Apr-1996 #text\_change 09-Jun-2000
C;Accession: S48654 #sequence\_revision 19-Apr-1996 #text\_change 09-Jun-2000
FRBS Lett. 352, 155-158, 1994
A;Hill, J.; Tyas, L.; Phylip, L.H.; Kay, J.; Dunn, B.M.; Berry, C.
FRBS Lett. 352, 185-188, 1994
A;Hille: High level expression and characterisation of Plasmepsin II, an aspartic protein A;Reference number: S48654
A;Reference number: S48654
A;Reference number: S48654
A;Reference number: Diamary
A;Reference number: Diamary
A;Residues: 1-20 <HIL>

Gapa ö Query Match 35.1%; Score 27; DB 2; Length 20; Best Local Similarity 50.0%; Pred. No. 4.2e+02; Matches 6; Conservative 1; Mismatches 5; Indels

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| | | :||| QMGRGSEHLTIG 20 3 ORGPGRAFVIIG 14 σ 요 ઠે

RESULT 3 PA0109

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dibydrolipoamide S-acetyltransferase (EC 2.3.1.12) - bovine (fragment)

C;Species: Bos primigenius taurus (cattle)
C;Species: Jo-Apr-1991 #sequence_revision 30-Apr-1991 #text_change 09-Jul-2004
C;Accession: A37823
R;Rahmatullah, M.; Radke, G.A.; Andrews, P.C.; Roche, T.E.
R;Rahmatullah, M.; Radke, G.A.; Andrews, P.C.; Roche, T.E.
A;Ritle: Changes in the core of the mammalian-pyruvate dehydrogenase complex upon selectifaccession: A37823; MUD:90354445; PMID:2167319
A;Recession: A37823
A;Roccession: A37823
A;Roccession: A37823
A;Roccession: A37823
exo-alpha-sialidase (EC 3.2.1.18) - influenza A virus (strain A/FPV/Rostock/34 [H7N1]) (1
                                  NyAlternate names: neuraminidase
C;Species: influenza A virus
C;Species: 1.Dec-1990 #sequence_revision 31-Dec-1990 #text_change 22-Jun-1999
C;Accession: 531286
R;Robertson, J.S.
NyIcleic Acids Res. 6, 3745-3757, 1979
A;Title: 5' and 3' terminal nucleoride sequences of the RNA genome segments of influenza A;Reference number: 511286, MUID:80034428; PMID:493121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nydrogensulfite reductase (EC 1.8.99.3) alpha chain - Desulfovibrio vulgaris (fragment)
Nydremate names: bisulfite reductase; desulfofuscidin; desulforubidin; desulforubidin;
C;Specias: Desulfovibrio vulgaris
C;Date: 19-Mar-1997 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004
C;Accession: S21197
C;Accession: S21197
Bur. J. Biochem. 205, 111-115, 1992
A;Title: The third subunit of desulfoviridin-type dissimilatory sulfite reductases.
A;Accession: S21197
A;Molecule type: protein
A;Accession: S21197
A;Molecule type: protein
A;Accession: S21197
A;Molecule type: protein
A;Csos-references: UNIPROT: P45574
A;Experimental source: strain Hildenborough
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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C;Complex: heterohexamer; two alpha, two beta and two gamma chains
C;Function:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            32.5%; Score 25; DB 2; Length 25; 36.4%; Pred. No. 1.2e+03; tive 4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 32.5%; Score 25; DB 2; Length 12; Best Local Similarity 44.4%; Pred. No. 5.9e+02; Matches 4; Conservative 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A,Map position: segment 6
C,Superfamily: influenza virus exo-alpha-sialidase
C,Keywords: glycosidase; hydrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 32.5
Best Local Similarity 36.4
Matches 4; Conservative
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porin porl - Arabidopsis thaliana (fragment)

porin porl - Arabidopsis thaliana (mouse-ear cress)

C;Spate: 30-Jun-1992 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004

C;Accession: PA0045

R;Kamo, M.; Kawakami, T.; Miyatake, N.; Tsugita, A.

submitted to JIPID, July 1994

A;Description: Separation and characterization of Arabidopsis proteins by two-dimensions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7
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porin por 1B - Arabidopsis thaliana (fragment)
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 07-Apr-1995 #sequence_revision 26-May-1995 #text_change 09-Jul-2004
C;Accession: PA0109
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Pred. No. 5e+02;
1; Mismatches 1; Indels
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Pred. No. 3.7e+02;
2; Mismatches 3;
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Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                       A, Cross-references: UNIPROT: QBLA84; UNIPROT: Q42292 A, Experimental source: root
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                                                                                                                                                          R;Kamo, M.; Kawakami, T.; Tsugita, A.submitted to JIPID, March 1995
A;Reference number: PA0109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A,Molecule type: protein
A,Residues: 1-14 «KAM»
A,Crose-references: UNIPROT:Q7MIW9
A,Experimental source: root
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Similarity 71.4%;
5; Conservative
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Best Local Similarity 50.0%;
Matches 6; Conservative
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A; Accession: PA0045
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Best Local Similarity
Matches 5; Conserv
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GRSFVPI 9
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RESULT 4

ઠે 셤 RESULT 6 S11286

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J. Exp. Med. 174, 1467-1476, 1991

A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental allergic A;Ederence number: PHO891; MUID:92078857; PMID:1836012

A;Accession: PHO915

A;Molecule type: mRNA

A;Residues: 1-14 <GOL>
A;Experimental source: concanavalin A-activated lymphoblast
A;Note: the authors translated the codon GGG for residue 8 as Glu and GAG for residue 9 c
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       dissimilatory sulfite reductase alpha chain, soluble - Desulfovibrio desulfuricans (fragm c)Species: Desulfovibrio desulfuricans (c)Species: Desulfovibrio desulfuricans (c)Species: Desulfovibrio desulfuricans (c)Species: Desulfovibrio desulfuricans (c)Species: Desulfovibrio desulfovibrio (s)Accession: S6490

R;Steuber, J; Arendsen, A.F.; Hagen, W.R.; Kroneck, P.M.H.

Eur. J. Biochem. 233, 873-879, 1995

A;Title: Molecular properties of the dissimilatory sulfite reductase from Desulfovibrio (A)Sterence number: S63489; MUID:96085152; PMID:8521853

A;Accession: S63490

A;Stetus: preliminary

A;Molecule type: protein

A;Residues: 1-20 <STE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          S31427
biliary glycoprotein - human
C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 07-Feb-1997
C;Accession: S31427
R;Nedellec, P.; Turbide, C.C; Barnett, T.R.; Beauchemin, N.
R;Nedellec, P.; Purbide, C.C; Barnett, T.R.; Beauchemin, N.
R;Nedellec, P.; Purbide, C.C; Barnett, T.R.; Beauchemin, N.
R;Nedellec, P.; Turbide, C.C; Barnett, T.R.; Beauchemin, N.
R;Nedellec, P.; Turbide, C.; Barnett, T.R.; Beauchemin, N.
R;Nedellec, P.; Turbide, C.; Barnett, T.R.; Beauchemin, N.
A;Nedellec, Interacterization of the human biliary glycoprotein regulatory region.
A;Status: preliminary
A;Nolecule type: DNA
A;Nesidues: 1-21 <NRE>
A;Cross-references: EMBL:X67277
C;Keywords: glycoprotein
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                                                                                                                                                                                                                                                  31.2%; Score 24; DB 2; Length 14;
50.0%; Pred. No. 1e+03;
cive 2; Mismatches 2; Indels
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llarity 36.4%; Pred. No. 1.4e+03;
Conservative 4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 24; DB 2; 1
Pred. No. 1.5e+03;
0; Mismatches 1;
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Best Local Similarity 50.00,
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Beet Local Similarity 80.00,
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Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                             T-cell receptor beta chain V-D-J region (100-4AE) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Accession: PT0515
R;Feeney, A.J.
Bxp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Note: sequence extracted from NCBI backbone (NCBIN:56069, NCBIP:56164)
C;Comment: This J3 segment may not be functional because of substitutions in the 7 mer
C;Keywords: heterotetramer; immunoglobulin
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C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Dacession: C2-May-1994 #sequence_revision 18-Nov-1994 #text_change 16-Aug-1996
C;Accession: C52375
R;Ayadi, H.; Marche, P.N.; Cazenave, P.A.
Immunogenetics 34, 201-207, 1991
A;Ttlle: Evolution of the rabbit immunoglobulin kappa chain genes.
A;Reference number: A53275; MUID:91372868; PMID:1909995
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C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
C;Accession: PH0915
R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
                                                                                                                                                        Gaps
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                                                                                                       Length 17;
                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 24; DB 2; Length 7; Pred. No. 2.8e+05; 1; Mismatches 0; Indels
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                                                                                                    Score 24.5; DB 2;
Pred. No. 9.9e+02;
1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: mRNA
A;Residues: 1-7 <FEE>
A;Experimental source: adult thymus, strain BALB/c
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
  A;Residues: 1-17 <RAH>
A;Cross-references: UNIPROT:Q7M2M8
C;Keywords: acyltransferase; coenzyme A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31.2%;
80.0%;
                                                                                                  Query Match
31.8%;
Best Local Similarity 66.7%;
Matches 6; Conservative
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Matches 4; Conservative
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Best Local Similarity
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A;Molecule type: DNA
A;Residues: 1-13 <AXA>
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GPGQA 7
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Cytochrome-c oxidase (EC 1.9.3.1) chain VII c, hepatic - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Bate: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: $65388; $65389
R;Schaegger, H.; Noack, H.; Halangk, W.; Brandt, U.; von Jagow, G.
Rix, J. Blochem. 230, 235-24, 1995
Bur, J. Blochem. 230, 235-24, 1995
A;Title: Cytochrome-c oxidase in developing rat heart. Enzymic properties and amino-termi
A;Reference number: $65388
A;Scatus: preliminary
A;Molecule type: protein
A;Residues: 1-10, eSCH>
A;Residues: 1-10, eSC
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C;Date: 03.Feb-1994 #sequence_revision 03-Feb-1994 #text_change 23-Mar-1995
C;Accession: PS0453
R;Tsugita, A;Pspid: April 1993
A;Reference number: PS0206
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30.5%; Score 23.5; DB 2;
Best Local Similarity 50.0%; Pred. No. 1.2e+03;
Matches 6; Conservative 1; Mismatches 4;
                                                                                                    RESULT 17
PS0453
36K protein 3124 - rice (etrain Nihonbare) (fragment)
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llarity 50.0%; Pred. No. 1.1e+03;
Conservative 2; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                   A,Molecule type: protein
A,Residues: 1-13 <TSU>
A,Experimental source: leaf, chlorophyll, stem
A,Note: molecular weight 36K, pl 6.1
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      1 GIGAAILSAGK 11
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Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                           A; Accession: PS0453
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          a
C;Species: Homo sapiens (man)
C;Date: 10-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 20-Jun-2000
C;Date: 10-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 20-Jun-2000
C;Datesion: C42866
R;Liu, Z.; Diaz, L.A.; Haas, A.L.; Giudice, G.J.
J. Biol. Chem. 267, 15829-15835, 1992
A;Title: CDNA cloning of a novel human ubiquitin carrier protein. An antigenic domain spthis human epidermal transcript.
A;Reference number: A42856; MUID:92348449; PMID:1379239
A;Accession: C42856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MSEL-neurophysin - African clawed frog (fragment)
N;Alternate names: vacopressin-associated neurophysin
C;Species: Xenopus laevis (African clawed frog)
C;Accession: B60422
R;Chacession: B60422
R;Chauvet, J.; Michel, G.; Rouille, Y.; Chauvet, M.T.; Acher, R.
R;Chauvet, J.; Michel, G.; Rouille, Y.; Chauvet, M.T.; Acher, R.
A;Title: Identification of two types of neurophysins in Xenopus laevis neurointermediate
A;Reference number: A60422; MUID:91067001; PMID:2250763
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bombinin-like peptide 4 - Bombina orientalis

bombinin-like peptide 4 - Bombina orientalis

C;Species: Bombina orientalis

C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 09-Jul-2004

C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 09-Jul-2004

C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 09-Jul-2004

S;Gibson, B.W.; Tang, D.; Mandrell, R.; Kelly, M.; Spindel, E.R.

J. Biol. Chem. 266, 23103-23111, 1991

A;Title: Bombinin-like peptides with antimicrobial activity from skin secretions of the A;Reference mumber: A41575; MUID:92078177; PMID:1744108
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                                                                                                                                                                                                                                                                                                             A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-22 -LIU5
A;Residues: 1-22 -LIU5
A;Experimental source: keratinocyte
A;Experimental source: keratinocyte
A;Note: sequence extracted from NCBI backbone (NCBIN:109895, NCBIP:109899)
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45.5%; Pred. No. 1.7e+03;
tive 2; Mismatches 4; Indels
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100.0%; Pred. No. 1.5e+03;
tive 0; Mismatches 0;
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A;Residues: 1-24 <CHA>
C;Superfamily: oxytocin-neurophysin
C;Keywords: pitultary
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C, Superfamily: bombinin H precursor
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Best Local Similarity 100.0
Matches 4; Conservative
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Matches 4; Conservative
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Matches 5, Conservative
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dentinal fluid transport-stimulating peptide - rat

NiAlternate names: DFT-stimulating peptide
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Species: DJML-2000 #sequence_revision 20-Jun-2000 #text_change 16-Aug-2004
C;Accession: JJ0001
R;Yamamoto, T.; Kobayashi, M.; Kobayashi, M.; Yamamoto, M.; Nomura, M.; Aonuma, S.
Cham. Pharm. Bull: 34, 3803-3811, 1986
A;Title: Isolation and amino acid sequence of dentinal fluid transport-stimulating peptic
A;Reference number: JJ0001; MUID:87131231; PMID:3815601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A,Molecule type: protein
A,Residues: 1-20 <YAM>
A,Cross-references: INIRROT:P07448
A,Experimental source: parotid gland
C,Comment: This peptide stimulates the transport of dentinal fluid, which is important for C,Keywords: hormone; parotid gland
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C;Superfamily: pyruvate dehydrogenase (lipoamide) alpha chain; thiamin pyrophosphate-bind
C;Keywords: oxidoreductase
                                                                       C;Species: Bos primigenius taurus (cattle)
C;Species: Bos primigenius taurus (cattle)
C;Aacession: 831220
R;Castillo, G.M.; Templeton, D.M.
R;Castillo, G.M.; Templeton, D.M.
A;Title: Subunit structure of bovine ESF (extracellular-matrix stabilizing factor(s)). A
A;Reference number: 831219; MUID:93178646; PMID:7680011
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Pred. No. 3.1e+03;
0; Mismatches 4; Indels
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100.0%; Pred. No. 3.1e+03;
tive 0; Mismatches 0;
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Best Local Similarity 55.6%;
Matches 5; Conservative
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A; Status: preliminary
A; Molecule type: protein
A; Residues: 1-20 <CAS>
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Best Local Similarity
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                                                  82K protein - bovine
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       A;Cross-references: UNIPROT:O52748; GB:BA000019; PIDN:BAB74000.1; PID:g17131393; GSPDB:C
A;Experimental source: strain PCC 7120
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Function:
A;Pathway: oxidative phosphorylation; respiratory chain
C;Keywords: electron transfer; membrane-associated complex; mitochondrial inner membrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Accession: S65629
R;Taketani, S.; Yoshinaga, T.; Furukawa, T.; Kohno, H.; Tokunaga, R.; Nishimura, K.; Ino R:Taketani, S.; Yoshinaga, T.; Furukawa, T.; Kohno, H.; Tokunaga, R.; Nishimura, K.; Ino A;Title. 230, 760-765, 1995
A;Title. Induction of terminal enzymes for heme biosynthesis during differentiation of m A;Reference number: S65629; MUID:95331315; PMID:7607249
A;Accession: S65629
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C;Superfamily: phytoene dehydrogenase
C;Keywords: heme blosynthesis; mitochondrion; oxidoreductase; porphyrin blosynthesis
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A;Reference number: S77980
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C;Species: Bos primigenius taurus (cattle)
C;Date: 14-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 26-May-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                 cytochrome-c oxidase (EC 1.9.3.1) chain VIII.1 - bigeye tuna (fragment) C;Species: Thurnus obesus (bigeye tuna) C;Date: 17-Sep.1997 #sequence_revision 17-Sep-1997 #text_change 09-Jul-2004 C;Accession: S77991
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                                                                                                                                               Query Match 29.9%; Score 23; DB 2; L
Best Local Similarity 66.7%; Pred. No. 1.8e+03;
Matches 4; Conservative 1; Mismatches 1;
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12 ERGSGR 17
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Best Local Similarity
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A;Residues: 1-17 <KUR>
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                                                                                                  A;Gene: patS
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Matches
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Jahuanawylase inhibitor - durum wheat (fragment)
Nahternate names: glutenin low molecular weight chain
Cispecies: Triticum durum (durum wheat)
Cipacies: Triticum durum (durum wheat)
Cipacession: 19-Mar-1997 #sequence_revision 29-Aug-1997 #text_change 09-Jul-2004
Cipacession: S10850
Rikobrehal, K.; Alary, R.
J. Sci. Food Agric. 48, 441-452, 1989
A;ritle: Isolation and partial characterisation of two low molecular weight durum wheat of the second recomber: S10849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytochrome-c oxidase (BC 1.9.3.1) chain VIIc - bigeye tuna (fragment)
C;Species: Thunnus obesus (bigeye tuna)
C;Species: 17-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 09-Jul-2004
C;Accession: 877990
E;Arcossion: 877990
E;Arnold, S.; Lee, J.; Kim, M.; Song, E.; Linder, D.; Lottspeich, F.; Kadenbach, B. submitted to the Protein Sequence Database, June 1997
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                                                                                           Score 22; DB 2; Length 25;
Pred. No. 3.8e+03;
0; Mismatches 5; Indels
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C;Superfamily: peroxidase
C;Keywords: blocked amino end; glycoprotein; oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pregnancy-specific glycoprotein SBU-3-62 - sheep (fragment)
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C;Superfamily: wheat alpha-amylase inhibitor
C;Keywords: alpha-amylase inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A,Cross-references: UNIPROT:Q9TRE1
C,Superfamily: Pepsin
C,Keywords: glycoprotein
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57.1%;
                                                                                                      Query Match 28.6%;
Best Local Similarity 44.4%;
Matches 4; Conservative
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Best Local Similarity 57.1
Matches 4; Conservative
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Best Local Similarity
Matches 5; Conserva
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S77990
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S2221
peroxidase (EC 1.11.1.7) - imperfect fungus (Arthromyces ramosus) (fragment)
peroxidase (EC 1.11.1.7) - imperfect fungus (Arthromyces ramosus)
C;Species: Arthromyces ramosus
C;Date: 12-Peb-1998 #sequence_revision 17-Apr-1998 #text_change 12-Jul-2004
C;Accession: S2221
R;Kjalke, M.; Andersen, M.B.; Schneider, P.; Christensen, B.; Schuelein, M.; Welinder, Biochim. Biophys. Acta 1120, 248-256, 1992
A,Title: Comparison of structure and activities of peroxidases from Coprinus cinereus, A,Title: Comparison of structure and activities of A,Recession: S22221
A,Recession: S22221
A;Molecule type: protein
A,Residues: 1-25 < KJA>
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C;Species: Homo sapiens (man)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 23-Jul-1999
C;Accession: S47N U · m---f.
                                                                                                                                                                                                                                                                                                 T-cell receptor beta chain (BTB15) - bovine (fragment)
C;Species: Bos primigenius taurus (cattle)
C;Date: 20-Peb-1995 #sequence_revision 20-Peb-1995 #text_change 20-Peb-1995
C;Accession: P00070
                                                                                                                                                                                                                                                                                                                                                                                                                                          R.Tanaka, A.; Ishiguro, N.; Shinagawa, M.
submitted to JIPID, May 1990
A.;Bescriptichen: Sequence analysis of bovine T-cell receptor beta chain genes.
A.;Reference number: U00472
A;Reference number: U00472
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A; Status: preliminary
A; Molecule type: mRNA
A; Residues: 1-23 < PLA.
A; Cross-references: EMBL: X71051; NID: 9506974; PIDN: CAA50368.1; PID: 9510653
C; Superfamily: immunoglobulin V region; immunoglobulin homology
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Pred. No. 3.5e+03;
1; Mismatches 5; Indels
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33.3%; Pred. No. 3.4e+03;
iive 2; Mismatches 4;
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Kriplaza, A.; Kono, D.H.; Theofilopoulos, A.N.
submitted to the EMBL Data Library, February 1993
A;Reference number: S40133
                     2; Mismatches
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Best Local Similarity 45.5%;
Matches 5; Conservative .
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Best Local Similarity 33.3
Matches 3; Conservative
                        3; Conservative
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A, Residues: 1-22 < TAN>
A, Experimental source: T cell
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C,Keywords: receptor
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Cyaccession: H41299
R;Cenatsu, Y.; Wege, H.; Straus, A.; Ott, M.; Bannwarth, W.; Lanchbury, J.; Panayi, G.; E
R;Cenatsu, Y.; Wege, H.; Straus, A.; Ott, M.; Bannwarth, W.; Lanchbury, J.; Panayi, G.; E
Proc. Natl. Acad. Sci. U.S.A. 89, 8534-8538, 1991
A;Title: The T-cell-receptor repertoire in the eynovial fluid of a patient with rheumatoi
A;Reference number: A41299; MUID:92020887; PMID:1656449
A;Recession: H41299
A;Status: preliminary
A;Residues: preliminary
A;Ross-references: GB:557504; NID:9236332; PIDN:AAB19963.1; PID:9236333
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: A42411
R;Leachman, S.A.; Gallagher, P.J.; Herring, B.P.; McPhaul, M.J.; Stull, J.T.
Aj. Biol. Chem. 267, 4930-4938, 1992
A;Title: Biochemical properties of chimeric skeletal and smooth muscle myosin light chair
A;Reference number: A42411; MUID:92165861; PMID:1371510
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A;Title: The molecular basis of homocystinuria due to cystathionine beta-synthase deficit A;Reference number: I51879; MUID:95282779; PMID:7762555
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   C;Species: Homo sapiens (man)
C;Date: 28-May-1992 #sequence_revision 28-May-1992 #text_change 05-Nov-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               myosin light chain kinase - chicken
C;Species: Gallus gallus (chicken)
C;Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cystathionine beta-synthase - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 02-Jul.1996 #sequence_revision 02-Jul-1996 #text_change 09-Jul-2004
C;Accession: IS1879
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80.0%; Pred. No. 3.8e+03;
tive 0; Mismatches 1; Indels
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A;Experimental source: skeletal muscle
A;Note: sequence extracted from NCBI backbone (NCBIP:84332)
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Pred. No. 3.8e+03;
1; Mismatches 4;
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A,Status: preliminary: translated from GB/EMBL/DDBJ
A,Molecule type: DIA
A,Residues: 1-16 <RES>
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Best Local Similarity 44.4%;
Matches 4; Conservative
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Best Local Similarity 80.0°
Matches 4; Conservative
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Matches 5, Conservative
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CiSpecies: Avena sativa (oat)
CiDate: 02-Dec193 #sequence_revision 27-Feb-1997 #text_change 09-Jul-2004
CiDate: 02-Dec1993 #sequence_revision 27-Feb-1997 #text_change 09-Jul-2004
CiAccession: S33800
R;Mummert, E.; Grimm, R.; Speth, V.; Eckerskorn, C.; Schiltz, E.; Gatenby, A.A.; Schaefe Nature 363, 644-648, 1993
A;Title: A TCP1-related molecular chaperone from plants refolds phytochrome to its photo A;Reference number: S33800; MUID:93288140; PMID:8099715
A;Accession: S33800
A;Accession: S33800
A;Accession: Liniary
A;Molecule type: protein
A;Residues: 1-13 <MUM>A;Accession: S3800
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A;Pathway: oxidative phosphorylation; respiratory chain
C;Keywords: electron transfer; membrane-associated complex; mitochondrial inner membrane
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C;Species: Jo.Gep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1147
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:99094761; PMID:1460419
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                                                                                                                                                                                                                                                                                                                                                      Query Match 27.3%; Score 21; DB 2; Length 10; Best Local Similarity 75.0%; Pred. No. 2.5e+03; Matches 3; Conservative 1; Mismatches 0; Indels
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A, Molecule trype: DNA
A, Residuces: 1-14 < WAS.
C, Keywords: heterotetramer; immunoglobulin
                                                    A,Molecule type: protein
A,Residues: 1-10 (ARN)
A,Cross-references: UNIPROT:P80982
A,Experimental source: heart; liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Cross-references: UNIPROT: Q7M1G8
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A;Reference number: $77980
A;Accession: $77990
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4, Conserv
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GPGK 9
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                                                                                                                                                                                                           A;Genome: nuclear
                                                                                                                                                                                   C, Genetica:
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26.6%;
ilarity 55.6%;
Conservative
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Matches 5, Conservative
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Matches 5; Conserv
17 GKGRRF 22
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A; Residues: 1-18 < LAN>
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                                                                          RESULT 39
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LX-1 tumor antigen - human (fragment)
LX-1 tumor antigen - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change 05-Jan-1996
C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change 05-Jan-1996
R;Rosenbaum, L.C.; Neuwelt, E.A.; Van Tol, H.H.M.; Loh, Y.P.; Verbalis, J.G.; Hellstroem Proc. Natl. Acad. Sci. U.S.A. 87, 9928-9932, 1990
Proc. Natl. Acad. Sci. U.S.A. 87, 9928-9932, 1990
A;Title: Expression of neurophysin-related precursor in cell membranes of a small-cell lay. A;Reference number: A39269; MUID:91088624; PMID:1702222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tex delta chain V-D-J region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Daccession: 149037
C;Accession: 149037
C;Accession: 22, 491-498, 1992
Eur. J. Immunol. 22, 491-498, 1992
A;Fittle: Mouse autoreactive gamma/delta T cells. II. Molecular characterization of the TA;Reference number: A49037; MUD:92164730; PMID:1311262
                                                                                                                                                     C;Accession: S09722
R;Monsalve, R.I.; Menendez-Arias, L.; Lopez-Otin, C.; Rodriguez, R.
ERS Lett. 263, 209-212, 139.
A;Title: beta-furns as structural motifs for the proteolytic processing of seed proteins
A;Reference number: S09720; MUID:90242974; PMID:2185951
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                                                                                                           C,Species: Brassica napus (rape)
C,Date: 19-Mar-1997 #sequence_revision 13-Mar-1998 #text_change 13-Mar-1998
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A,Molecule type: DNA
A,Residues: 1-19 < EEZQ.
A,Cross-references: GB:S90660; NID:g246304; PIDN:AAB21555.1; PID:g246305
A,Cross-reference: dendritic epidermal T-cell lines
A,Note: sequence extracted from NCBI backbone (NCBIN:90660, NCBIP:90671)
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Pred. No. 5e+03;
0; Mismatches 2; Indels
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50.0%; Pred. No. 4.4e+03;
tive 1; Mismatches 4;
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57.1%; Pred. No. 4.2e+03;
ive 1; Mismatches 2;
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A;Residues: 1-22 <ROS>
C;Superfamily: oxytocin-neurophysin
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Best Local Similarity 66.7%;
Matches 4; Conservative
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Matches 4; Conservative
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A; Molecule type: protein
A; Residues: 1-9;10-18 «MON>
A; Experimental source: seed
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GGGRIWRLIG 11
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6 RIPKQPG 12
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A; Molecule type: DNA
A; Residues: 1-18 - SBN >
A; Crose-refered GB: M13798; NID: 9204707; PIDN: AAA41371.1; PID: 9554447
A; Crose-refered GB: M13798; NID: 9204707; PIDN: AAA41371.1; PID: 955447
B; Lang, P.; Mocikat, R.
Gene 102, 261-264, 1991
A; Title: Immunoglobulin heavy-chain joining genes in the rat: comparison with mouse and PA; Reference number: JH0666; MUID: 91340162; PMID: 1908401
A; Accession: JH0666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C, Accession: A25941; JH0666
R; Brueggemann, M.; Free, J.; Diamond, A.; Howard, J.; Cobbold, S.; Waldmann, H. Proc. Natl. Acad. Sci. US.A. 83; 6075-6079, 1986
A; Pitle: Immunoglobulin heavy chain locus of the rat: striking homology to mouse antibody A; Reference number: A25941; MUID:86287397; PMID:3016742
                                                                                                                                                                                                                            endogenous type C retrovirue
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NiAlternate names: T-kinin
Cippecies: Homo sapiens (man)
Cipte: 02-Dec-1993 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
Biol. Chem. Hoppe-Seyler 371, 977-981, 1990
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probable gas polymerase pseudogene - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 13-Aug-1999 #sequence_revision 13-Aug-1999 #text_change 20-Oct-2000
C;Date: 17-Aug-1999 #sequence_revision 13-Aug-1999 #text_change 20-Oct-2000
C;Date: 17-Aug-1999 #sequence_revision 18.Repaske, R.; O'Neill, R.R.; Steele, P.E.; Martin, M.A.
Proc. Natl. Acad. Sci. U.S.A. 80, 678-682, 1983
A;Title: Characterization and partial nucleotide sequence of endogenous type
A;Reference number: Z14423; MUID:83143994; PMID:6298769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ig heavy chain J-H1 region - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 05-Jun-1988 #sequence_revision 05-Jun-1988 #text_change 23-Jul-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 24;
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33.3%; Pred. No. 5.4e+03;
ive 3; Mismatches 7;
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Pred. No. 5.1e+03;
1; Mismatches 2;
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A;Cross-references: EMBL:J00274; NID:g182154
C;Keywords: pseudogene
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Gaps

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C;Date: 20-Jul-1990 #sequence_revision 20-Jul-1990 #text_change 09-Jul-2004
C;Accession: C34874
R;McMahon, G.; Davis, B.F.; Huber, L.J.; Kim, Y.; Wogan, G.N.
Proc. Natl. Acad. Sci. U.S.A. 87, 1104-1108, 1990
A;Title: Characterization of c-Ki-ras and N-ras oncogenes in aflatoxin B-1-induced rat li
A;Reference number: A34874; MUID:90138946; PMID:2105496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T. Cell receptor alpha chain V region (clone 2PBL V alpha 24-6) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1799
C;Accession: PH1799
C;Accession: PH179, 2. Y Yockey, C.E.; Brenner, M.B.; Balk, S.P.
J. Exp. Med. 178, 1-16, 1993
A;Title: Analysis of T cell antigen receptor (TCR) expression by human peripheral blood (
                                                                                                                                                                                                                                                                                                                Score 20; DB 2; Length 15;
Pred. No. 5.3e+03;
3; Mismatches 2; Indels
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Pred. No. 5.6e+03;
1; Mismatches 0
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                                                                                                                                                                                      A,Status: preliminary
A,Molecule type: protein
A,Residues: 1-15 <MCM-
A,Cross-references: UNIPROT:Q7M030
                                                                                                                                                                                                                                                                                                                h 26.0%;
Similarity 44.4%;
4; Conservative 3
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Best Local Similarity 75.0%;
Matches 3; Conservative
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Best Local Similarity
Matches 4; Conserv
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A; Status: preliminary
A; Molecule type: mRNA
A; Residues: 1-16 < POR>
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PH1790
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Bur. J. Biochem. 221, 1111-1116, 1994

Bur. J. Biochem. 221, 1111-1116, 1994

A;Title: Jachtification of tissue-specific isoforms for subunits Vb and VIIa of cytochro A;Reference number: 543624; MUID:94237150; PMID:8181469

A;Accession: S43634
  A;Title: Ile-Ser-bradykinin is an aberrant permeability factor in various human malignar
A;Reference number: $13279; MUID:91166748; PMID:2076202
A;Accession: $13279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Genetics:
A;Genome: nuclear
C;Keywords: cardiac muscle; heart; membrane-associated complex; mitochondrion; oxidoredu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    constant region genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytochrome-c oxidase (EC 1.9.3.1) chain VIIc, cardiac - rainbow trout (fragment) C;Species: Oncorhynchus mykiss (rainbow trout) C;Date: 20-Oct-1994 #sequence_revision 01-Nov-1996 #text_change 16-Jul-1999
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A;Molecule type: DNA
A;Residues: 1-15 - KOY>
A;Cross-references: GB:M14159; NID:g338852; PIDN:AAA60679.1; PID:g553690
C;Keywords: T-cell receptor
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R;Toyonaga, B.; Yoshikai, Y.; Vadasz, V.; Chin, B.; Mak, T.W.
Proc. Natl. Acad. Sci. U.S.A. 82, 8624-8628, 1985.
A;Title: Organization and sequences of the diversity, joining, and A;Reference number: A94081; MUID:86094276; PMID:3866244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Homo sapiens (man)
C;Date: 16-Aug-1988 #sequence_revision 16-Aug-1988 #text_change
                                                                                                                                                                                                                    Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26.0%; Score 20; DB 2; I
75.0%; Pred. No. 5.3e+03;
tive 1; Mismatches 0;
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                                                                                                                                                                                                                 Score 20; DB 2;
Pred. No. 4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: protein
A;Residues: 1-15 <FRE>
A;Note: the source is designated as Salmo gairdneri
                                                                                                                                                                                                                                                                     0; Mismatches
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C;Species: Rattus norvegicus (Norway rat)
                                                                                 A;Molecule type: protein
A;Residues: 1-11 wWN>
A;Cross-references: UNIPROT:Q7M4P1
C;Keywords: bradykinin
                                                                                                                                                                                                                 26.0%;
                                                                                                                                                                                                              Query Match 26.0
Best Local Similarity 66.7
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 75.0 Matches 3; Conservative
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us-08-869-386-3.rapb

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May 16, 2005, 12:57:08 ; Search time 113.231 Seconds (without alignments) 70.804 Million cell updates/sec
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2: /cgn2 6/ptodata/1/pubpaa/PCT NEW PUB.pep:*
3: /cgn2 6/ptodata/1/pubpaa/US06 NEW PUB.pep:*
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13: /cgn2 6/ptodata/1/pubpaa/US09 NEW PUB.pep:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries

    protein search, using sw model

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Gapop 10.0 , Gapext 0.5
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

	Description	Sequence 160, App	Sequence 25, Appl	Sequence 3, Appli	Sequence 3, Appli	Sequence 24, Appl	Sequence 1, Appli	Sequence 1, Appli	Sequence 45, Appl	Sequence 15, Appl	Sequence 24, Appl	Sequence 8, Appli	Sequence 9, Appli	Sequence 3, Appli
	QI		US-10-178-488-25	US-09-813-659-3	US-10-283-610A-3	US-09-827-345-24	US-10-311-111-1	US-10-398-932-1	US-10-062-710-45	US-09-810-310-15	US-09-810-310-24	US-09-989-621-8	US-09-827-688-9	US-09-077-439A-3
	8	17	14	σ	15	10	14	16	14	Q	0	σ	10	20
	Query Match Length DB	24	21	20	20	20	20	20	18	15	15	15	15	15
æ	Query Match	93.5	76.4	74.8	74.8	73.2	71.5	71.5	66.7	62.6	62.6	62.6	62.6	62.6
	Score	115	94	92	92	90	88	88	85	77	77	77	77	77
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Sequence 246, App	Sequence 202, Appli Sequence 6, Appli	7	Sequence 44, Appl	Sequence 186, App	113	154	536	Sequence 6, Appli	155	21, 7	112	Sequence 114, App	12, 7	535	7	σÌ	15,	Sequence 50, Appl	Ä	Sequence 1, Appli	22,	40,	Sequence 3, Appli	'n	Sequence 10, Appl	Sequence 158, App	12, 7	533,	Seguence 12, Appl	Sequence 24, Appl
7.	4 US-10-133-210-262  4 US-10-147-910-6	2	14 US-10-062-710-44	14 US-10-239-313A-186		10-621-	14 US-10-239-313A-536	17 US-10-622-003-6	17 US-10-621-675-155	10 US-09-993-307-21	15 US-10-373-592-112	IS US-10-373-592-114	17 US-10-628-004-12	14 US-10-239-313A-535	17 US-10-621-675-7	(7 US-10-621-675-9	LO US-09-956-940-15	10 US-09-956-940-50	LS US-10-360-647A-1	17 US-10-613-018-1	17 US-10-613-018-22	17 US-10-613-018-40	14 US-10-311-111-3	16 US-10-398-932-3	US-09-901-106-10	17 US-10-621-675-158	17 US-10-621-675-12	14 US-10-239-313A-533	US-09-901-106-12	4 US-10-178-488-24
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14	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	. 45

## ALIGNMENTS

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MERGUT.

US.-10-621-675-160

Sequence 160, Application US/10621675

Sequence 160, Application US/10621675

Sequence 160, Application US/10601639841

SEQUENCE 160, Application US/1080163084

SEQUENCE 160, Application US/1080163084

SEQUENCE 160, USCOSO04939841

SEQUENCE 160, USCOSO04939841

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING

TITLE OF INVENTION: PRIVATE PEPTIDES CORRESPONDING THEM THEM TO INMUNOLOGICALLY IMPORTANT

TITLE OF INVENTION: BETTORES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS

TITLE OF INVENTION: CONTAINING THEM

FILE REPERENCE 2752-11

FRICK APPLICATION NUMBER: US/00/516, 834A

PRIOR PELING DATE: 1996-09-10

PRIOR PELING DATE: 1993-03-06

NUMBER OF SEQ ID NOS: 600

SOFTWARE: PARIOR TOTAL ON WHERE: EPT/2093/00517

PRIOR PELING DATE: 1993-03-06

NUMBER OF SEQ ID NOS: 600

SOFTWARE: PARIOR TOTAL ON WHERE: EPT/2093/00517

PRIOR PELING DATE: 1993-03-06

NUMBER OF SEQ ID NOS: 600

SOFTWARE: PARIOR THEM

SEQ ID NO 160

SORGANISM: Human immunodeficiency virus

US-10-61-675-160

QUERT WARCH

BEST CORSETVATIVE ON : Mismatches 1; Indels 0; Gaps 0;
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Query Match
Best Local Similarity 95.0
Matches 19; Conservative
                                                         19; Conservative
          Query Match
Best Local Similarity
Matches 19; Conserv
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US-09-827-345-24
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; Sequence 3. Application US/09813659
; Patent No. US20020012989A1
; GENERAL INFORMATION:
; APPLICANT: Ledbetter, Jeffrey A.
; APPLICANT: Linsley, Peter S.
; APPLICANT: Bajorath, Jurgen
APPLICANT: Billiland, Lisa K.
; APPLICANT: Billiland, Lisa K.
; APPLICANT: Gilliland, Lisa K.
; APPLICANT: Gilliland, Lisa K.
; TITLE OF INVENTION: EXPERSION OF PRODUCING BISPECIFIC FUSION PROTEINS
; TITLE OF INVENTION: EVISION PROTEINS IN A MAMMALIAN CELL
; TITLE OF INVENTION: PUSION PROTEINS IN A MAMMALIAN CELL
; TITLE OF INVENTION: By 194502
; TITLE OF INVENTION: WINBER: US/09/813,659
; CURRENT FILING DATE: 2000-04-13
; PRIOR FILING DATE: 1993-04-13
; PRIOR FILING DATE: 1993-09-13
; PRIOR FILING DATE: 1993-02-01
; NUMBER OF SEQ ID NOS: 32
; SOFTWARE: PATENTIN VEY: 2.0
                                                                                                                                      Sequence 25, Application US/10178488
; Publication No. US20030165535A1
; APPLICANT: Rovinski, Benjamin
; APPLICANT: Rao, Fed-Long
; APPLICANT: Yao, Fed-Long
; APPLICANT: Yao, Fed-Long
; APPLICANT: Wichel N. Michel H.
; TITLE OF INVENTION: PLUBALITY OF MUTATIONS
; FILE REFERENCE: 1038-1238 MIS.
; FILE REFERENCE: 1038-1238 MIS.
; CURRENT APPLICATION NUMBER: US/10/178,488
; RIOR FILING DATE: 202-10-25
; RRIOR FILING DATE: 1999-02-26
; NUMBER OF SEQ ID NOS: 26
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 76.4%; Score 94; DB 14; Length 21; Best Local Similarity 90.5%; Pred. No. 3.9e-07; Matches 19; Conservative 0; Mismatches 2; Indels
1 NNTRKSERIORGPGRAFVTIGKIG 24
                          2 NTRKSERIQRGPGRAFVTIGK 22
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ORGANISM: Unknown Organism
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                                                                                                                           RESULT 2
US-10-178-488-25
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publication No. US20030021800A1

publication No. US20030021800A1

publication No. US20030021800A1

general information:

APPLICANT: CHERMANN, JEAN-CLAUDE

APPLICANT: CHERMANN, JEAN-CLAUDE

APPLICANT: GALEA, PASCALE

TITLE OF INVENTION: VACCINA PREVENTION FOR THE TREATMENT AND

TITLE OF INVENTION: INTRACELULAR PHASE, COMPOSITION FOR THE TREATMENT AND

TITLE OF INVENTION: PREVENTION OF HIV INFECTIONS, ANTIBODIES AND METHOD OF

TITLE OF INVENTION: DAGNOSIS

TITLE OF INVENTION: DAGNOSIS

TITLE OF INVENTION: DAGNOSIS

TITLE OF INVENTION: OLSO-1-0216

CURRENT APPLICATION NUMBER: US/09/827,345

CURRENT APPLICATION NUMBER: 09/599,549

PRIOR FILING DATE: 2000-06-23

PRIOR FILING DATE: 1996-06-28

PRIOR FILING DATE: 1996-06-28

PRIOR APPLICATION NUMBER: 08/973,551

PRIOR FILING DATE: 1998-02-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAPPLICANT: Ledbetter, Jeffrey A.

APPLICANT: Hayden, Martha S.

APPLICANT: Hayden, Martha S.

APPLICANT: Hayden, Martha S.

APPLICANT: Hayden, Martha S.

APPLICANT: Linaley, Perer S.

APPLICANT: Bajorath, Jurgen

APPLICANT: Brilland, Lisa K.

TITLE OF INVENTION: ENGINE WORTON WECTORS ENCODING BISPECIFIC FUSION PROTEINS

TITLE OF INVENTION: FUSION PROTEINS IN AMMALIAN CELL

TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC

TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC

TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC

TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC

FILE REPERENCE: ONLOTE/30436.18053, 436

PRIOR FILING DATE: 2000-04-13

PRIOR APPLICATION NUMBER: 08/121,054

PRIOR APPLICATION NUMBER: 08/121,054

PRIOR PRIOR APPLICATION NUMBER: 08/121,054

PRIOR PRIOR PRING DATE: 1993-09-13

PRIOR PRING DATE: 1993-09-13

PRIOR APPLICATION NUMBER: 08/13,420

PRIOR FILING DATE: 1993-02-01

NUMBER OF SEQ ID NOS: 35

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 3

LENGTH: 20

MARCHARY DATE: 1093-02-01

SEQ ID NO 3
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Length 20;
                                                                 i; Indels
74.8%; Score 92; DB 9; I 95.0%; Pred. No. 7.4e-07; iive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 3, Application US/10283610A
; Publication No. US20030219876A1
; GENERAL INFORMATION:
                                                                                                                                                                                      1 RKSIRIQRGPGRAFVTIGKI 20
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PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetically Designed
OS-10-398-932-1
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GENERAL INFORMATION:
APPLICANT: Khleif, Samir N.
TITLE OF INVENTION: IMMUNOLOGICAL RESPONSES TO PEPTIDE ANTIGENS; FILE REFERENCE: 15280-415100US; CURRENT APPLICATION NUMBER: US/09/810,310; CURRENT APPLICATION NUMBER: 05/189,396; PRIOR APPLICATION NUMBER: 60/189,396; PRIOR PILING DATE: 2000-03-15; NUMBER OF SEQ ID NOS: 61; SOFTWARE: Patentin Ver. 2.1; SEQ ID NO 15; LENGTH: 15
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APPLICANT: Li, Frank Q.

APPLICANT: Chu, Yong-Liang
APPLICANT: Oth, Jian-Tai

TITLE OF INVENTION: Polymeric Conjugates for Delivery of
TITLE OF INVENTION: MHC-Recognized Epitopes
TITLE OF INVENTION: Wia Peptide Vaccines
FILE REFERENCE: 3781-001-27

CURRENT FILING DATE: 2002-02-05

PRIOR PELICATION NUMBER: US 60/310,498

PRIOR PELICATION NUMBER: US 60/310,498

PRIOR PELICATION NUMBER: US 60/310,498

PRIOR PILING DATE: 2001-08-08

NUMBER OF SEQ ID NOS: 232

SOFTWARE: FRASEQ for Windows Version 4.0

SEQ ID NO 45

LENGTH: 18
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100.0%; Pred. No. 2.1e-05;
tive 0; Mismatches 0;
                                                                                                                                                    Query Match 71.5%; Score 88; DB 16; Best Local Similarity 90.0%; Pred. No. 3e-06; Matches 18; Conservative 0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , OTHER INFORMATION: HIV Helper-T Cell Epitopes
US-10-062-710-45
                                                                                                                                                                                                                                                                                                                                                                              US-10-062-710-45

"Sequence 45, Application US/10062710
; Publication No. US20030049253A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                       4 RKSERIQRGPGRAFVTIGKI 23
    TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 16; Conserva
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Publication No. US20030121065A1

GENERAL INFORMATION:
APPLICANT: SHIBA, KIYOTAKA
TITLE OF INVENTION: MULTIFUNCTIONAL BASE SEQUENCE AND ARTIFICIAL GENE CONTAINING THE FILE REFERENCE: 4439-4004
CURRENT APPLICATION NUMBER: US/10/311,111
PRIOR PILING DATE: 2002-12-13
PRIOR FILING DATE: 2000-16097
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 34
SEQ ID NO 1
IENGTH: 20
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Sequence 1, Application US/10398932

Publication No. US20040171803A1

SERERAL INFORMATION:

APPLICANT: SHIBA, KIYOTAKA

APPLICANT: OHNO, TSUNEYA

TITLE OF INVENTION: OF EPITOPE

TITLE OF INVENTION: OF EPITOPE

TITLE OF INVENTION: OF EPITOPE

CURRENT APPLICATION NUMBER: US/10/398,932

CURRENT FILING DATE: 2003-04-11

PRIOR FILING DATE: 2003-04-10-10

PRIOR PELICATION NUMBER: JP 2000/314288

PRIOR PELING DATE: 2000-10-10

PRIOR APPLICATION NUMBER: JP 2000/314288

RHOR PELING DATE: 2001-0-10

SOFTWARE: PATENTIN VOT: 2.1

SSOFTWARE: PATENTIN VOT: 2.1
                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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Pred. No. 1.5e-06;
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94.7%; Pred. No. 1...
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PRIOR APPLICATION NUMBER: FR 95/07914
PRIOR FILING DATE: 1995-06-30
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 24
LENGTH: 20
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; OTHER INFORMATION: Designed peptide
US-10-311-111-1
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                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
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Matches 18; Conservative
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Best Local Similarity
Matches 18; Conserv
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US-10-311-111-1
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                                                                                                                                         TYPE: PRT
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Fublication No. US2003020289A1

GENERAL INFORMATION:

APPLICANT: Collier, R. John

APPLICANT: Blanke, Steven R.

APPLICANT: Banson, Enicka L.

APPLICANT: Banson, Michael N.

TITLE OF INVENTION: Handles for Delivering Compounds into Cells

FILE REFERENCE: 00246/187002

CURRENT APPLICATION NUMBER: US/09/077,439A

CURRENT APPLICATION NUMBER: US/09/077,439A

CURRENT APPLICATION NUMBER: US/09/077,439A

CURRENT FILING DATE: 1996-12-13

FRIOR FILING DATE: 1996-12-13

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FRIOR FILING DATE: 1995-12-13

FRIOR FILING DATE: 1995-12-13
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                          62.6%; Score 77; DB 9; Length 15; llarity 100.0%; Pred. No. 0.0001; Conservative 0; Mismatches 0; Indels
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ORGANISM: Homo sapien
                                     Query Match
Best Local Similarity
Matches 15; Conserva
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; ORGANISM: HIV p18
US-09-827-688-9
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Sequence 24, Application US/09810310

Patent No. US20020044948A1

GENERAL INFORMATION:

APPLICANT: Khielf, Samir N.

APPLICANT: Khielf, Samir N.

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR CO-STIMULATION OF TITLE OF INVENTION: IMMUNOLOGICAL RESPONSES TO PEPTIDE ANTIGENS

FILE REFERENCE: 15280-415100US

CURRENT APPLICATION NUMBER: 2001-03-14

PRIOR APPLICATION NUMBER: 60/189,396

PRIOR APPLICATION NUMBER: 60/189,396

NUMBER OF SEQ ID NOS: 61

SOFTWARE: Patentin Ver. 2.1
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; Sequence 8, Application US/09989621
; Patent No. US20020151683A1
; Patent No. US20020151683A1
; GREERAL INFORMATION:
   APPLICANT: Mogam Biotechnology Research Institute
   APPLICANT: Chang, Jin-Soo
   APPLICANT: Chang, Jin-Soo
   APPLICANT: Chang, Jin-Soo
   APPLICANT: Choi, Sung-Yoo
   APPLICANT: Choi, Myeong
   APPLICANT: 
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                                                                                                     Query Match 62.6%; Score 77; DB 9; Length 15; Best Local Similarity 100.0%; Pred. No. 0.0001; Matches 15; Conservative 0; Mismatches 0; Indels
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SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
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; OTHER INFORMATION: ANTIGEN US-09-810-310-15
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LENGTH: 15
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| Sequence 2, Application US/10787880
| Sequence 2, Application WS-102050025777A1
| Sequence 2, Application Wo. US20050025777A1
| Publication No. US20050025777A1
| APPLICANT: Sheehy, Michael J.
| PULE REFERENCE: 033267-018
| PILE REFERENCE: 033267-018
| CURRENT APPLICATION NUMBER: US/10/787,880
| CURRENT PILIG DATE: 2004-02-27
| PRIOR PLICATION NUMBER: US/09/191,772
| PRIOR PLICATION NUMBER: US/09/191,772
| NUMBER OF SEQ ID NOS: 2
| SOFTWARE: FastSEQ for Windows Version 4.0
| SEQ ID NO 2: LENGTH: 15
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100.0%; Pred. No. 0.0001;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                     APPLICANT: Fuller, Deborah
APPLICANT: Fuller, James
APPLICANT: Fuller, James
APPLICANT: Shipley, Timothy
TITLE OF INVENTION: Vaccine Composition
FILE REFERENCE: 03257-006
CURRENT APPLICATION NUMBER: US 60/291,654
PRIOR PILING DATE: 2001-05-18
PRIOR PILING DATE: 2001-05-18
PRIOR PILING DATE: 2001-05-18
PRIOR PILING DATE: 2001-05-18
NUMBER OF SEQ ID NOS: 53
SOFTWARE: PESTSEQ for Windows Version 4.0
SEQ ID NO 6
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US-10-062-710-44
; Sequence 44, Application US/10062710
                                                                                                                                                                                                                                 Sequence 6, Application US/10147910; Publication No. US20030124718A1; GENERAL INFORMATION:
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Best Local Similarity 100.0%
Matches 15; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
8 RIQRGPGRAFVTIGK
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US-10-787-880-2
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; ORGANISM: HIV
US-10-147-910-6
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                                                                                                                                                                                                                                                                                                                          Sequence 246, Application US/10133210

Sequence 246, Application US/10133210

Publication No. US20030103964A1

GENERAL INFORMATION:
APPLICANT: Defisi, Charles
APPLICANT: Berzofeky, Jay
APPLICANT: Waccaro, Dennis
APPLICANT: Waccaro, Dennis
APPLICANT: Wang, Chao
TITLE OF INVENTION: COMPOSITIONS THEREOF
TITLE OF INVENTION: COMPOSITIONS THEREOF
FILE REPERENCE: BU-035AX
CURRENT APPLICATION NUMBER: US/10/133,210
CURRENT PILING DAIE: 2002-04-26

MUMBER OF SEQ ID NOS: 281

SOFTWARE: PATCHIN Ver. 2.0

SEQ ID NO 246

LENGTH: 15
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Sequence 262, Application US/10133210

Publication No. US20030103964A1

SEQUENCE TO USEQUENCE

APPLICANT: DeLisi, Charles

APPLICANT: Berzofeky, Jay

APPLICANT: Galukota, Kamalakar

APPLICANT: Vaccaro, Dennis

APPLICANT: Weng, Zhiping

APPLICANT: Convertion: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND

TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND

TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND

TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND

TITLE OF INVENTION: WETHODS FOR DESIGNING MOLECULAR CONJUGATES AND

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; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-10-133-210-262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-10-133-210-246
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 62.6%; Score 77; DB 14; Length 15; Best Local Similarity 100.0%; Pred. No. 0.0001; Matches 15; Conservative 0; Mismatches 0; Indels
   Best Local Similarity 100.0%; Pred. No. 0.0001;
Matches 15; Conservative 0; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
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                                                                                                                                                1 RIQRGPGRAFVTIGK 15
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                                                                                                       8 RIORGPGRAFVTIGK 22
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US-10-133-210-246
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
APPLICANT: HAYNES, BARTON F.
APPLICANT: KORBER, BETTE T.
APPLICANT: DE LORIMIER, ROBERT M.
TITLE OF INVENTION: POLYVALENT INMUNOGEN
FILE REFERENCE: 1579-785
CURRENT APPLICATION NUMBER: US/10/373,592
CURRENT FILING DATE: 2003-012-26
PRIOR PILING DATE: 2002-11-07
PRIOR PILING DATE: 2002-11-07
PRIOR PILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 120
SOFTWARE: PALENTING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 120
SOFTWARE: PALENTING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 120
SOFTWARE: PALENTING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 120
SOFTWARE: PALENTING DATE: 2001-11-07
SOFTWARE: PALENTING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 120
SOFTWARE: PALENTING DATE: 2001-11-07
SOFTWARE: PALENTING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 120
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                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 76.2
Matches 16; Conservative
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US-10-239-313A-186

Sequence 186, Application US/10239313A

Publication No. US20030175285A1

GENERAL INFORMATION:

APPLICANT: KLINGUER - HAMOUR, Christine

APPLICANT: CORVAIA, Nathalie

APPLICANT: GOETSCH, Liliane

TITLE OF INVENTION: N-TERMINAL A GLUTAMINE IN THE FORM

TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID

TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID

TITLE OF INVENTION: NUMBER: US/10/239,313A

CURRENT FILING DATE: 2002-09-19

PRIOR PILING DATE: 2000-03-23

PRIOR PLING DATE: 2000-03-22

NUMBER OF SEQ ID NOS: 697

SOFTWARE: PATENTIN VET. 2.1

SEQ ID NO 186

LENGTH: 15

LENGTH: 15
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100.0%; Pred. No. 0.00011;
7ative 0; Mismatches 0; Indels
                              GENERAL INFORMATION:
APPLICANT: Li, Frank Q.
APPLICANT: Chu, Yong-Liang
APPLICANT: Oth, Yong-Liang
APPLICANT: Oth, Jian-Tai
TITLE OF INVENTION: Polymeric Conjugates for Delivery of
TITLE OF INVENTION: Was Peptide Waccines
FILE REFERENCE: 3781-001-27
CURRENT FILING DATE: 2002-05
FRIOR APPLICATION NUMBER: US 60/310,498
FRIOR FILING DATE: 2001-08-08
NUMBER OF SEQ ID NOS: 232
SOFTWARE: FastSEQ for Windows Version 4.0
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Best Local Similarity 100.0%; Pred. No. 0.00057;
Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: HIV Helper-T Cell Epitopes US-10-062-710-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
          Publication No. US20030049253A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 RIORGPGRAFVTIGK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 RIORGPGRAFVTIGK 22
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Best Local Similarity
Matches 15; Conserv
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Gaps

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56.1%; Score 69; DB 15; Length 22; 76.2%; Pred. No. 0.0024; tive 0; Mismatches 3; Indels

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### PEPLICANI: DG LAFY, KODESTC

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING

TITLE OF INVENTION: TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR USE IN

TITLE OF INVENTION: BIOTINYLATED PEPTIDES CORRESPONDING THEM ANTICOLOGICALLY IMPORTANT

TITLE OF INVENTION: CONTAINING THEM

FILE OF INVENTION: CONTAINING THEM

FRICH REPERRICE: 2752-11

CURRENT APPLICATION NUMBER: US/09/576, 824A

FRIOR APPLICATION NUMBER: 09/146,028

FRIOR PILING DATE: 1993-11-22

FRIOR FILING DATE: 1993-03-06

FRIOR FI
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; ORGANISM: Human immunodeficiency virus
US-10-621-675-154
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| NNTRKS--IHIGPGRAFYTTGEI 21
                                                                                                                                                                                                                                                                                        ; Sequence 154, Application US/10621675; Publication No. US20050049398A1; GENERAL INFORMATION:
4 NNTRKS--IQIGPGRAFYTTG 22
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Matches 16; Conserv
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RESULT 22

; Sequence 113, Application US/10373592; Publication No. US20040001851A1; GENERAL INFORMATION:

US-10-373-592-113

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RESULT 26
US-10-373-592-112
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                                                                    GENERAL INFOGRATION:
APPLICANT: CCRVAIA, Nathalie
APPLICANT: CCRVAIA, Nathalie
APPLICANT: CCRVAIA, Nathalie
APPLICANT: CCRVAIA, Nathalie
TAPLICANT: BEEK, Alain
APPLICANT: GOSTSCH, Liliane
TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMIC IN THE FORM
TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID
FILE REFERENCE: 343 727 - US
CURRENT APPLICATION WHBER: US 100-03-19
FRIOR FILLING DATE: 2000-03-19
PRIOR FILLING DATE: 2000-03-23
PRIOR FILLING DATE: 2001-03-22
NUMBER OF SEQ ID NOS: 697
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 5:6
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PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR USE IN
A PROCESS FOR DETERMINATION OF ANTIBODIES OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gapa
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; Sequence 6, Application US/10622003
; Publication No. US20050014230A1
; GENERAL INFORMATION: The Company of FULLY HUMAN ANTIBODIES
; TILLE REFERENCE: 16865-002001
; CURRENT APPLICATION NUMBER: US/10/622,003
; CURRENT FILING DATE: 2003-07-16
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: FASESEQ for Windows Version 4.0
; SEQ ID NO 6
; SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 55.3%; Score 68; DB 14; Length 13; Best Local Similarity 100.0%; Pred. No. 0.002; Matches 13; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Synthetically generated peptide US-10-622-003-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Human immunodeficiency virus
Sequence 536, Application US/10239313A Publication No. US20030175285A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 155, Application US/10621675
Publication No. US20050049398A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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TITLE OF INVENTION: P
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-239-313A-536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-621-675-155
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BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
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Pred. No. 0.0046;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: HAYNES, Joel R.
APPLICANT: HAYNES, Joel R.
TITLE OF INVERTINGYN, JOSHUA
TITLE OF INVERTINGYN, WUCLEIC ACID ADJUVANTS
FILE REFERENCE: APF41.20
CURRENT APPLICATION NUMBER: US/09/993,307
CURRENT APPLICATION NUMBER: 60/253,381
PRIOR APPLICATION NUMBER: 60/253,381
PRIOR FILING DATE: 2000-11-27
NUMBER OF SEQ ID NOS: 26
SEQ ID NO 21
TITLE OF INVENTION: BIOTINYLATED PEFTIDES CO TITLE OF INVENTION: EPITOPES, A PROCESS FOR TITLE OF INVENTION: CONTAINING THEM FILE REPERENCE: 2752-11

CURRENT APPLICATION NUMBER: US/10/621,675
CURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: US/09/576,824A

PRIOR APPLICATION NUMBER: 09/123,425
PRIOR PILING DATE: 1996-09-30

PRIOR PILING DATE: 1996-09-30

PRIOR PILING DATE: 1993-11-22

PRIOR PILING DATE: 1993-0-68

PRIOR PILING DATE: 1993-0-68

PRIOR PILING DATE: 1993-0-68

PRIOR PILING DATE: 1993-0-68

PRIOR PILING DATE: 1993-0-69

PRIOR PILING DATE: 1993-0-69

PRIOR PILING DATE: 1993-0-69

SOFTWARE: PATENTION NUMBER: EP 92400598.6

NUMBER OF SEQ ID NOS: 600

SOFTWARE: PATENTING NOS: 600

SOFTWARE: PATENTING NOS: 600
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Publication No. US20040001851A1
GENERAL INFORMATION:
APPLICANT: HAYNES, BARTON F.
APPLICANT: KORBER, BETTE T.
APPLICANT: DE LORIMIER, ROBERT M.
TITLE OF INVENTION: POLYVALENT IMMUNGEN
FILE REFERENCE: 1579-785
CURRENT APPLICATION NUMBER: US/10/373,592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 NNTRKSERIQRGPGRAFVTIGKI 23
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US-09-993-307-21
Sequence 21, Application US/09993307
Publication No. US20030162733A1
GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               53.7%;
86.7%;
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1 Similarity 69.6%;
16; Conservative
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; ORGANISM: synthetic construct
US-09-993-307-21
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Best Local Similarity
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Best Local Similarity
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ACTIVITY AGAINST PRIMARY HIV-1 STRAINS
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Best Local Similarity
                       TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                        FEATURE:
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| Sequence 12, Application US/10628004
| Publication No. US20050058983A1
| GENERAL INFORMATION:
| APPLICANT: ABGENIX, INC.
| APPLICANT: PUBLIC HEALTH RESEARCH INSTITUTE
| APPLICANT: HE, VUXIAN
| APPLICANT: HE, VUXIAN
| APPLICANT: CORVALAN, USE RR.
| APPLICANT: OF INVENTION: USE OF TRANSGENIC MICE FOR THE EFFICIENT ISOLATION OF TITLE OF INVENTION: NOVEL HUMAN MONOCLONAL ANTIBODIES WITH NEUTRALIZING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                FEATURE:
7. OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-10-373-592-112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: Synthetic prince of the Information: peptide US-10-373-592-114
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52.0%; Score 64; DB 15; Length 22;
Best Local Similarity 71.4%; Pred. No. 0.014;
Matches 15; Conservative 0; Mismatches 4; Indels
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52.0%; Score 64; DB 15;
Best Local Similarity 71.4%; Pred. No. 0.014;
Matches 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: HAXINES, BARTON F.
APPLICANT: KORBER, BETTE T.
APPLICANT: NCRBER, BETTE T.
TITLE OF INVENTION: POLYVALENT IMMUNGEN
FILE REFERENCE: 1579-785
CURRENT APPLICATION NUMBER: US/10/373,592
CURRENT FILING DATE: 2003-02-26
PRIOR APPLICATION NUMBER: 10/289,228
PRIOR PELING DATE: 2002-11-07
PRIOR PELING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 120
SOFTWARE: PATENTIN VEY: 2.1
PRIOR APPLICATION NUMBER: 10/289,228
PRIOR APPLICATION NUMBER: 10/289,228
PRIOR FILING DATE: 2002-11-07
PRIOR PLING DATE: 2001-11-07
PRIOR PLING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 120
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 112
LENGTH: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-373-592-114
; Sequence 114, Application US/10373592
; Publication No. US20040001851A1
; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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LENGTH: 22
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APPLICANT: De Leye, Robert

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
TITLE OF INVENTION: TO IMMUNOLOGICALLY IMPORTANT PETTOPES AND THEIR USE IN
TITLE OF INVENTION: A PROCESS FOR DETERMINATION OF ANTIBODIES OF
TITLE OF INVENTION: BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
TITLE OF INVENTION: EPITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
TITLE OF INVENTION: CONTAINING THEM
FILE REFERENCE: 2752-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 535, Application US/10239313A

Publication No. US20030175285A1

Publication No. US20030175285A1

GENERAL INFORMATION:

APPLICANT: CORVAIA, Nathalie

APPLICANT: CORVAIA, Nathalie

APPLICANT: BECK, Alain

TITLE OF INVENTION: NOTERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE FORM

TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID

FILE REFERENCE: 343 727 - US

CURRENT APPLICATION NUMBER: US/10/239,313A

CURRENT PILING DATE: 2002-09-19

FRIOR APPLICATION NUMBER: PCT 01/7072

PRIOR APPLICATION UNMER: DO 03-23

PRIOR APPLICATION UNMER: DO 03-22

NUMBER OF SEQ ID NOS: 697

SOFTWARE: PatentIN Ver. 2.1
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                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Synthetic; CTHER INFORMATION: peptide
US-10-628-004-12
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Pred. No. 0.01;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 NKRKRIHIQRGPGRAFYTTKNI 23
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FILE REFERENCE: ABX-PHRI PCT
CURRENT APPLICATION NUMBER: US/10/1
CURRENT FILING DATE: 2003-07-25
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 12
LENGTH: 24
                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 63.6%;
Matches 14; Conservative
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                                                                                              Score 60.5; DB 17; Length 24;
Pred. No. 0.052;
1; Mismatches 3; Indels 3
                                                                                                                                                                                                                                                                                                                                                                                                            ESSUL: 5-5-40-15
US-09-956-940-15
Sequence 15, Application US/09956940
Sequence 15, Application US/09956940
GENERAL INFORMATION:
APPLICANT: HAYNES, BARTON F.
TITLE OF INVENTION: USE OF SYNTHETIC PEPTIDES TO INDUCE
TITLE OF INVENTION: USE OF SYNTHETIC T AND B CELI
AUTOANTIGENS OR INFECTIOUS AGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: ARLINGTON
COUNTRY: U.S.A.

COUNTRY: U.S.A.

ZIP: 22201-4714

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRINT APPLICATION DATA:
APPLICATION DATE: US/09/956,940
FILING DATE: 12-Oct-2001
CLASSIFFCATION: «Unknown»
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48.0%; Score 59; DB 10; Length 13;
Best Local Similarity 92.3%; Pred. No. 0.045;
Matches 12; Conservative 0; Mismatches 1; Indels
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APPLICATION NUMBER: US/08/460,673

FILING DATE: JULKNOWN;

PILING DATE: 10-FEB-1993

APPLICATION NUMBER: US 07/833,429

FILING DATE: 10-FEB-1992

APPLICATION NUMBER: US 07/591,109

FILING DATE: 01-00-1990

APPLICATION NUMBER: US 07/591,109

FILING DATE: 08-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: NIXON & VANDERHYE P.C. STREET: 1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 1579-5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
;
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
US-09-956-940-15
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REGISTRATION NUMBER: 32,955
                                                                                                                                                                                                                                           1 NNTRKSERIORGPGRAFVTIGKI 23
                                                                                                                                                                                                                                                                      2 NNTRKS--IYIGPGRAFTT-GRI 21
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, OTHER INFORMATION: modified site US-10-621-675-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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INFORMATION FOR SEQ ID NO: 15:
                                                                                                           Query Match
Best Local Similarity 69.6%;
Matches 16; Conservative
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CORRESPONDENCE ADDRESS:
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APPLICANT: De Leys, Robert
TITLE OF INVENTION:
TITLE
TITL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Pred. No. 0.043;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Indels
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      CURRENT APPLICATION NUMBER: US/10/621,675
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US/09/576,824A
PRIOR APPLICATION NUMBER: 08/1723,425
PRIOR FILING DATE: 1996-09-30
PRIOR FILING DATE: 1993-011-22
PRIOR FILING DATE: 1993-01-122
PRIOR FILING DATE: 1993-03-08
PRIOR FILING DATE: 1993-03-06
NUMBER OF SEQ ID NOS: 600
SOFTWARE: PAECHIN VOICE: 1992-03-06
NUMBER OF SEQ ID NOS: 600
SOFTWARE: PAECHIN VET: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 NNTRKSERIORGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 9, Application US/10621675 Publication No. US20050049398A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: VARIANT
LOCATION: (24)
OTHER INFORMATION: modified site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: VARIANT
LOCATION: (1)
OTHER INFORMATION: modified site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: modified site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               49.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: VARIANT
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LOCATION: (24)
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8

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| SERERAL INFORMATION:
| JERREAL INFORMATION:
| APPLICANT: WEINHUES, URSULA-HENRIKE
| APPLICANT: HOSS, EVA
| APPLICANT: SEIDEL, CHISTOPH
| APPLICANT: SEIDEL, CHISTOPH
| APPLICANT: WIEDMANN, MICHAEL,
| TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING
| TITLE OF INVENTION: MILLIPIE ANTIGENS
| FILE REPERENCE: 100564-07003
| CURRENT FILING DATE: 100564-07003
| CURRENT FILING DATE: 1995-07-24
| PRIOR FILING DATE: 1994-07-25
| PRIOR PLING DATE: 1994-07-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: Epitope region ; OTHER INFORMATION: of HIV type 1, HIV type 2 or HIV subtype 0 US-10-613-018-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 48.0%; Score 59; DB 17; Length 17; Best Local Similarity 73.7%; Pred. No. 0.06; Matches 14; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3; Indels
                                    APPLICANT: HOSS, Eva
TITLE OF INVENTION: METAL CHELATE-LABELLED PEPTIDES
FILE REFERENCE: 2923-529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 59; DB 15;
Pred. No. 0.06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: polypeptide/antigen/epitope
US-10-360-647A-1
                                                                                                                       CURRENT APPLICATION NUMBER: US/10/360,647A
CURRENT FILING DATE: 2003-02-10
PRIOR APPLICATION NUMBER: US 08/776189
PRIOR FILING DATE: 1997-01.24
PRIOR PILING DATE: 1995-07-24
PRIOR PLILING DATE: 1995-07-24
PRIOR PILING DATE: 1994-08-31
PRIOR PILING DATE: 1994-08-31
PRIOR FILING DATE: 1994-08-31
PRIOR FILING DATE: 1994-07-25
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin version 3.2
LENGTH: 17
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      Wienhues, Ursula-Henrike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/10613018 Publication No. US20050074750A1 GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity 73.7%;
Matches 14; Conservative
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                                                                                         RESULT 33
US-09-366-940-50
18-09-366-940-50
Sequence 50, Application US/09956940
Publication No. US20030022826A1
GENERAL INFORMATION:
APPLICANT: HAYNES, BARTON F
TITLE OF INVENTION: USE OF SYNTHETIC PEPTIDES TO INDUCE
TITLE OF INVENTION: USE OF SYNTHETIC PEPTIDES TO AUTOPES OF
AUTOANTIGENS OR INFECTIOUS AGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READBALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/956,940
FILING DATE: 12-Oct-2001
CLASSIFICATION: -Unknown>
PRIOR APPLICATION OF THE COMPANIENT APPLICATION NUMBER: US 08/015,987
FILING DATE: 10-FEB-1992
APPLICATION NUMBER: US 07/833,429
FILING DATE: 10-FEB-1992
APPLICATION NUMBER: US 07/833,429
FILING DATE: 10-FEB-1992
APPLICATION NUMBER: US 07/833,429
FILING DATE: 10-FEB-1990
APPLICATION NUMBER: US 07/93,854
FILING DATE: 01-OCT-1990
APPLICATION NUMBER: US 07/93,854
FILING DATE: 01-OCT-1990
APPLICATION NUMBER: US 07/093,854
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48.0%; Score 59; DB 10; Length 15;
Best Local Similarity 92.3%; Pred. No. 0.053;
Matches 12; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: WILSON, MARY J.
REGIGTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 1579-5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 50:
US-09-956-940-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/10360647A Publication No. US20040039178A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (703) 816-4100
TELEX: 200797 NIXN UR
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                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
ADDRESSEE: NIXON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Siedel, Christoph
1 TRKSIRIQRGPGR 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 TRKSERIQRGPGR 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: VIRGINIA
COUNTRY: U.S.A.
ZIP: 22201-4714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-360-647A-1
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Gaps

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Gaps 7

Length 17;

us-08-869-386-3.rapb

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US-10-311-111-3
Sequence 3) Application US/10311111
Publication No. US20030121065A1
Publication No. US20030121065A1
GENERAL INFORMATION:
APPLICANT: SHIBA, KIYOTAKA
TILLE OF INVENTION:
TILLE OF INVENTION:
CURRENT APPLICATION NUMBER: US/10/311,111
CURRENT APPLICATION NUMBER: 2002-12-13
PRIOR APPLICATION NUMBER: JP 2000-180997
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PATENTIN VETBION 3.1
SOFTWARE: PATENTIN VETBION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic peptide
TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING
TITLE OF INVENTION: MULTIPLE ANTIGENS
FILE REPERENCE: 100564-07003
CURRENT APPLICATION NUMBER: 1201-07
FIRIOR APPLICATION NUMBER: PCT/EP95/02919
FRIOR PLICATION NUMBER: PCT/EP95/02919
FRIOR FILING DATE: 1995-07-24
FRIOR FILING DATE: 1995-07-25
FRIOR PLING DATE: 1995-07-25
FRIOR PLING DATE: 1994-08-31
FRIOR FILING DATE: 1094-08-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 59; DB 17; Length 21;
Pred. No. 0.075;
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Pred. No. 0.064;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (3) OTHER INFORMATION: Epsilon-aminocaproic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 NNTRKSERIQRGPGRAFVT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 NNTRKS--ISIGPGRAFYT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTHER INFORMATION: Beta-alanine US-10-613-018-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (2) OTHER INFORMATION: Beta-alanine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial Sequence
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Best Local Similarity 91.7%;
Matches 11; Conservative
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NAME/KEY: MOD_RES
LOCATION: (1)
OTHER INFORMATION: BPRu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: MOD_RES
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NAME/KEY: MOD_RES
LOCATION: (4)
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                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: SEIDEL, CHRISTOPH
APPLICANT: WIEDMANN MICHAEL
TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING
TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING
TITLE OF INVENTION: WULTIPLE ANTIGENS
FILE REFERENCE: 100564-07003
CURRENT PEPLICATION NUMBER: US/10/613,018
CURRENT PELLING DATE: 2003-07-07
PRIOR PILING DATE: 1995-07-24
PRIOR FILING DATE: 1995-07-24
PRIOR FILING DATE: 1994-07-25
PRIOR FILING DATE: 1994-07-25
PRIOR FILING DATE: 1994-09-31
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PALENTIN VET. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Pred. No. 0.075;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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PUBLICATION NO. US20050074750A1

PUBLICATION NO. US20050074750A1

APPLICANT: WEINHUES, URSULA-HENRIKE

APPLICANT: WEINSE-MULLER, CORNELIA

APPLICANT: PASTZ, ELKE

APPLICANT: PASTZ, ELKE

APPLICANT: SEIDEL, CHRISTOPH

APPLICANT: SEIDEL, CHRISTOPH

APPLICANT: WIEDMANN, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: MOD RES
LOCATION: (1)
OTHER INFORMATION: Digoxigenin-3-cme
                                                                                                                                                                                                                                                                       APPLICANT: WEINHUES, URSULA-HENRIKE
                                                                                                                                                                                                                                                                                                                                      HOSS, EVA
FAATZ, ELKE
OFENLOCH-HAHNLE, BEATUS
                                                                                                                                                                                                                                                                                                        KRUSE-MULLER, CORNELIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 NNTRKSERIORGPGRAFVT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 NNTRKS--ISIGPGRAFYT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; LOCATION: (4) _
; OTHER INFORMATION: Beta-alanine
US-10-613-018-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (2) OTHER INFORMATION: Beta-alanine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: MOD_RES
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Gaps

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Gaps .. 0

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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Synthetically Designed OTHER INFORMATION: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                      ## Sequence 3, Application US/10398932

| Sequence 3, Application US/10398932
| Publication No. US20040171803A1
| GENERAL INFORMATION
| APPLICANT: SHIBA, KIYOTAKA
| APPLICANT: SHIBA, KIYOTAKA
| TITLE OF INVENTION: APTIFICIAL PROTEINS WITH ENRICHED IMMUNOGEN
| TITLE OF INVENTION: OF EPITOPE
| FILE REFERENCE: 024918-0103
| CURRENT FILING DATE: 2003-04-11
| PRIOR FILING DATE: 2003-04-11
| PRIOR FILING DATE: 2001-10-10
| PRIOR FILING DATE: 2001-10-10
| PRIOR FILING DATE: 2000-10-13
| NUMBER OF SEQ ID NOS: 52
| SOFTWARE: Patentin Ver: 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPER ELOPPY disk

COMPUTER: INDEP C compatible

COMPUTER: INDEP C compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Ralease #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/901,106

FILING DATE: 10-Jul-2001

CLASSIFCATION: «UNKNOWN»

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 58; DB 16; Length 13;
Pred. No. 0.064;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 10, Application US/09901106
Patent No. US20020151667A1
GENERAL INFORMATION:
APPLICANT: Garoff, Henrik
Liljestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based on Alphaviruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/07/920,281C
FILING DATE: 13-AUG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Murphy Jr., Gerald M.
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 828-103P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      47.2%;
91.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
2 RIORGPGRIFVT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11; Conservative
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Best Local Similarity
Matches 11; Conserv
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                                                                                                         US-10-398-932-3
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GENERAL INTOJULIAN:

APPLICANT: De Leys, Robert

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF WITHING IN THEIR USE IN

TITLE OF INVENTION: A PROCESS FOR DETERMINATION OF ANTHEODIES OF

TITLE OF INVENTION: BUTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLGGICALLY IMPORTANT

TITLE OF INVENTION: BUTINYLATED PEPTIDES FOR PREPARING THEM AND COMPOSITIONS

TITLE OF INVENTION: CONTAINING THEM

TITLE OF INVENTION: CONTAINING THEM

FILE OF INVENTION: CONTAINING THEM

FRICH REPERRICE: 2752-11

CURRENT APPLICATION NUMBER: US/09/576, 824A

PRIOR APPLICATION NUMBER: 08/146,028

PRIOR PILING DATE: 1993-11-22

PRIOR FILING DATE: 1993-11-22

PRIOR FILING DATE: 1993-03-06

SOFTWARE: PATCHAIN VET. 2.1

SEQ ID NO 158

TENNAME.
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                                                                                                                                                                                                                                                                                                                                                        Score 58; DB 9; Length 15;
Pred. No. 0.075;
1; Mismatches 1; Indels
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                                                                                                                                                                                          TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 10:
US-09-901-106-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ch
1 Similarity 56.5%; Pred. No. 0
13; Conservative 2; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 NNTRKSERIQRGPGRAFVTIGKI 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |||||| | :||||
| NNTRKS--ITKGPGRVIYATGQI 21
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 12, Application US/10621675; Publication No. US20050049398A1; GENERAL INFORMATION:
                                                                             INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids
                                                      703-241-2848
                                                                                                                                                                                                                                                                                                                                                           47.28;
                                                                                                                                                                                                                                                                                                                                                                                     84.68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8 RIQRGPGRAFVTI 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 RIQRGPGRAFVEL 15
                                                                                                                                                                                                                                                                                                                                                                                                                 11; Conservative
                            TELEPHONE:
TELEFAX: 70
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: De Leys, R
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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Matches 13; Conserv
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GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-621-675-158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-621-675-158
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RESULT 44
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APPLICANT: CORVAIA, Nathalie
APPLICANT: GORVAIA, Nathalie
APPLICANT: GORVAIA, Nathalie
APPLICANT: GORYAIA, Nathalie
TITLE OF INVENTION: WOLECTLE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE FORM
TITLE OF INVENTION: OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION NUMBER: US/10/239,313A
CURRENT APPLICATION NUMBER: PR 00/03711
PRIOR APPLICATION NUMBER: PCT 01/70772
PRIOR FILING DATE: 2000-03-22
PRIOR FILING DATE: 2000-03-22
NUMBER OF SEQ ID NOS: 697
         EPITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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47.2%; Score 58; DB 17; Length 25;
Best Local Similarity 56.5%; Pred. No. 0.13;
Matches 13; Conservative 2; Mismatches 6; Indels
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100.0%; Pred. No. 0.076;
tive 0; Mismatches 0; Indels
TITLE OF INVENTION: EPITOPES, A PROCESS FOR TITLE OF INVENTION: CONTAINING THEM FILE OF INVENTION: CONTAINING THEM FILE OF INVENTION: CONTAINING THEM CURRENT APPLICATION NUMBER: US/10/621,675; CURRENT FILING DATE: 2003-07-18; PRIOR APPLICATION NUMBER: 08/723,425; PRIOR FILING DATE: 1996-09-30; PRIOR FILING DATE: 1996-09-30; PRIOR FILING DATE: 1993-03-08; PRIOR FILING DATE: 1923-03-08; PRIOR FILING DA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Mismatches
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Sequence 533, Application US/10239313A

PUBLIcation No. US20030175285A1

GENERAL INFORMATION:
APPLICANT: KLINGUER - HAMOUR, Christine
APPLICANT: CORVAIA, Nathalie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 NNTRKSERIORGPGRAFVTIGKI 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; LOCATION: (25)
; OTHER INFORMATION: modified site
US-10-621-675-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 11; Conserv
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OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: VARIANT
LOCATION: (25)
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Sequence 124, Application US/10178488
Sequence 124, Application US/10178488
Publication No. US2003016555A1
GENERAL INFORMATION:
APPLICANT: Rovinski, Benjamin
APPLICANT: Cao, Shi.xian
APPLICANT: You's Feilong
APPLICANT: Klein, Michel H.
TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES MADE NON-INFECTIOUS BY A
TITLE OF INVENTION: PURPALITY OF MUTATIONS
FILE REFERENCE: 1038-1238 MIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gape
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                                                                                                                                                                                                                                                                                            STATE: Virginia

COUNTRY: USA

ZIP: 22040-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/901,106
FILING DATE: 10-Jul-2001
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46.3%; Score 57; DB 9; Length 17; 100.0%; Pred. No. 0.12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                    APPLICANT: Garoff, Henrik
Liljestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based on
Alphaviruses
                                                                                                                                                                             NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/920,281C
FILING DATE: 13-AUG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Murphy Jr., Gerald M.
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 828-103P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-241-1300
TELEFAX: 703-241-2848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: linear; MOLECULE TYPE: protein; SEQUENCE DESCRIPTION: SEQ ID NO: 12: US-09-901-106-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/178,488 CURRENT FILING DATE: 2002-10-25
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              Sequence 12, Application US/09901106
Patent No. US20020151067A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 17 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11; Conservative
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Matches 11; Conserva
JS-09-901-106-12
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; PRIOR FILING DATE: 1999-02-26
; NUMBER OF SEQ ID NOS: 26
; SCOTYMARE: Patentin Ver. 2.1
; SEQ ID NO 24
; LENGTH: 19
; TYPE: PRT
; ORGANISM: Unknown Organism
; PRATURE:
; OTHER INFORMATION: Description of Unknown Organism: Artificial US-10-178-488-24
                                                                                                                                                                                                                                                                                                                 Query Match 46.3%; Score 57; DB 14; Length 19; Best Local Similarity 66.7%; Pred. No. 0.14; Matches 14; Conservative 1; Mismatches 4; Indels
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Search completed: May 16, 2005, 13:10:22 Job time: 113.231 secs g

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1;

Gaps

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; TELEPHONE: (212) 302-8989; TELEPHONE: (212) 302-8998; INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 24 amino acids; TYPE: amino acid; TOPOLOGY: circular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Appli
Sequence 6, Appli
Sequence 160, App
Sequence 110, App
Sequence 12, Appl
Sequence 13, Appl
Sequence 99, Appl
                                                                    May 16, 2005, 12:52:18 ; Search time 36.9231 Seconds (without alignments) 48.522 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Description
                                                                                                                                                                                                                                                                                                                                         Issued_Patents AA:*
1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
5.1.6
Compugen Ltd.
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US-08-109-148-6

US-08-123-425A-160

US-08-131-2

US-09-112-206-160

US-09-112-206-160

US-09-190-497A-160

US-09-50-497A-160

US-09-50-497A-160

US-09-50-497A-160

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US-08-447-506-13

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US-08-467-515-13

US-08-467-515-13

US-08-467-518-99

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US-08-465-217A-99

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US-08-462-507A-99
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version 5
                                                                                                                   US-08-869-386-3
123
1 NNTRKSERIQRGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                         Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                              protein search, using sw model
                                                                                                                                                                   BLOSUM62
Gapop 10.0 , Gapext 0.5
GenCore
(c) 1993
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                                                                                                                                                                                                                                                     seq length: 0 seq length: 25
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Match
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                                                                                                                     Title:
Perfect score:
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                                              OM protein
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Maximum DB
                                                                                                                                             Sequence:
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		(080977)  Intra  orizio  cina
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O 4 C - C Ω Ω 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4		Application 552766 FORMATION: T: DeRoseil T: DeRoseil T: Panozzo, T: Panozzo, T: Disello,
88 883 883 883 883 76 76 76 76 76 76 76 76		pplic Despective of the control of t
O O O O O O O O O O O O O O O O O O O		T 1  -097-751-1  -097-751-1  CHORNATION: APPLICANT: DEROSSH, APPLICANT: DEROSSH, APPLICANT: PRIOZZO, APPLICANT: PRIOZZO, APPLICANT: PRIOZZO, APPLICANT: DIBELLO, APPLICANTION: APPLICANTON: APPLICANTON SERVEN SON SERVEN SERVEN SON SERVEN SERVEN SON SERVEN SERVEN SON SERVEN SERVEN SERVEN SON SERVEN SERVEN SERVEN SON SERVEN SERVEN SERVEN SERVEN SON SERVEN
11111 00000000000000000000000000000000		T 1 -097-751-1 -097-751-1 ence L No. 554 ent No. 554 ent No. 557 APPLICANT: COUNTRY: ZITE OF II MEDIUM TY COMPUTER IO COMPUTER OPERATING SOFTWARE CURRENT APPLICAT: FLING DA CLASSIFIC FRING DA APPLICAT: FLING DA APPLICATION APPLI
		T I 1 097-75 one to be compared to b
00000000000000000000000000000000000000		SULT 1 -08-097-751-1 Sequence 1, Applicati Sequence 1, Applicati Sequence 1, Applicati APPLICANT: Descoss APPLICANT: Panciz APPLICANT: Panciz APPLICANT: Panciz APPLICANT: Panciz APPLICANT: Panciz APPLICANT: DiBell ADDRESSES: Hedm STREET: 1185 AV CITY: New York STREET: 10036 COMPUTER: New York STATE: New York STATE: New York COUNTRY: USA ZIP: 10036 COMPUTER: IBM P OPERATING SYSTEM BELICATION NUMB FILING DATE: 19 CLASSIFICATION NUMB FILING DATE: 19 CLASSIFICATION NUMB FILING DATE: 19 CLASSIFICATION NUMB FILING DATE: 10 APPLICATION NUMB FILING DATE: 10 CASSIFICATION NUMB FILING DATE: 10 REFERRNCE/DOCKET TELECOMMUNICATION REFERRNCE/DOCKET TELECOMMUNICATION
		RESULT 1 105-08-097-751-1 5 Sequence 1, A 7 Patent No. 55 GENERAL INFO APPLICANT: COMPUTER P MEDIUM T COMPUTER P MEDIUM T COMPUTER P APPLICAT APPLI

Length 24;

Score 115; DB 1; Pred. No. 2.5e-10;

93.5%; 95.8%;

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PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
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CORRESPONDENCE ADDRESS:
ADDRESSED.
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                       NUMBER OF SEQUENCES: 453
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/O8/146,028
INFORMATION FOR SEQ ID NO: 160:
SEQUENCE GHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                   Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/723,425A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93.5%; Score 115; DB 3;
95.8%; Pred. No. 2.5e-10;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 93.5%; Score 115; DB 2; Best Local Similarity 95.8%; Pred. No. 2.5e-10; Matches 23; Conservative 0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: NIXON & VANDERHYE, P.C.
STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
CITY: Arlington
STATE: VA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24
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; Sequence 160, Application US/08723425A
; Patent No. 6165730
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NAME: SADOFF, B.J.
REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 1487
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703.816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 160: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: DELEYS, ROBERT
TITLE OF INVENTION: PEPTIDE
TITLE OF INVENTION: PEPTIDE
TITLE OF INVENTION: ANTIBOD
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Best Local Similarity 95.8
Matches 23; Conservative
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MOLECULE TYPE: peptide
US-08-723-425A-160
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                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: peptide
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       FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: Biz
                                                                                                                                                                                                                                                                                   amino acid
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                                                                                                                                                                                                                                                                                                                                                                            US-08-146-028-160
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CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR
IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED
PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES,
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           IndelB
                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Mactico, Robert Allan
APPLICANT: Stockley, Peter George
APPLICANT: Talbot, Simon John
TITLE OF INVENTION: Antigen-Presenting Capsid with
TITLE OF INVENTION: Fusion MS2-Coat Protein
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
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Best Local Similarity 95.8%; Pred. No. 2.5%
Matches 23; Conservative 0; Mismatches
           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:

ZIP: 10022-2585

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5", 1.44MD
COMPUTER: IBM PS2-486
COMPUTER: OBJECTION DATA:
APPLICATION NUMBER: GB 9101550.3
FILING DATE: 01/24/91
APPLICATION NUMBER: PCT/GB92/00124
FILING DATE: 01/24/91
ATTORNEY AGENT INFORMATION:
NAME: NISSENDAUM, ISTAEL
REGISTRATION NUMBER: 27,582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 NNTRKSERIORGPGRAFVTIGKIG 24
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Patent No. 5891640
GENERAL INFORMATION:
                                                                                                                                                                                                                       Sequence 6, Application US/08090148 Patent No. 5534257
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Rosenman & Colin
STREET: 575 Madison Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 940-8636
TELEFAX: (212) 940-6404
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NOT RELEVANT
           23; Conservative
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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US-08-146-028-160
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US-08-090-148-6
                Matches
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                                                                                                                                                        Sequence 2, Application US/08480332

Batent No. 6180134

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Zalipsky, Samuel; Woodle, Martin; Martin, Francis;
APPLICANT: Barenholz, Yechezkel
ITILE OF INVENTION: Enhanced Circulation Effector Composition and
TITLE OF INVENTION: Method
ITILE OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SUGTWARE: PAPLICATION DATA:
APPLICATION NUMBER: US/08/480,332
FLING DATE: 7-UNN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/316,436
FLING DATE: 29-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/316,436
FLING DATE: 23-ABA-1993
ATTORNEY/AGENT INFORMATION:
NAME: MOIN JUMS W.
REGISTRATION NUMBER: 38,563
REFERENCE/DOCKET NUMBER: 53.25-0115.31
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Avenue, Suite 250
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide 2, Fig. 13
24
                        1 NNTRKSIRIQRGPGRAFVTIGKIG 24
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NNTRKSERIORGPGRAFVTIGKIG
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TELEFAX: (415) 324-0960
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 95.8<sup>3</sup>
Matches 23, Conservative
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INDIVIDUAL ISOLATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                RESULT 5
US-08-480-332-2
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; Sequence 160, Application US/09112206

RESULT 6 US-09-112-206-160

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CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTIVILATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES, PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
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PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
PROCESS FOR THE DETERMINATION OF ANTIBODIES OF
A PROCESS FOR DETERMINATION OF ANTIBODIES OF
BIOTINYTATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
EPITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
CONTAINING THEM
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Pred. No. 2.5e-10;
0; Mismatches 1.
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Pred. No. 2.5e-10;
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TITLE OF INVENTION: PROCESS FOR THE TITLE OF INVENTION: CORRESPONDING TO TITLE OF INVENTION: IN A PROCESS FOR TITLE OF INVENTION: PROFESS FOR PREPAINTE OF INVENTION: PROFESS FOR PREPAINTER OF SEQUENCES; 453

COMPUTER READABLE FORM:
MEDIUM TYPE: FLORPY disk
COMPUTER: IBM PC COMPAISH
COMPUTER: IBM PC COMPUTER: IBM 
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/146,028
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95.8%;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
TYPE: amino acid
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SEQ ID NO 14
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Best Local Similarity
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Best Local Simil
Matches 23; (
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PROCESS FOR THE DETERMINATION OF PEPTIDES
CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR
IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED
PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES,
PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
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PCT-US92-06688-12
; Sequence 12, Application PC/TUS9206688
; GENERAL INFORMATION
; APPLICANT: REPLIGEN CORPORATION
; APPLICANT: THE ROCKEFELLER UNIVERSITY
; TITLE OF INVENTION: WULTIPLE ANTIGEN PEPTIDES FOR USE AS HIV
; TITLE OF INVENTION: VACCINES
; NUMBER OF SEQUENCES: 21
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Archardson
STREET: 225 Franklin Street
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Pred. No. 2.5e-10;
0; Mismatches 1; Indels
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Pred. No. 2.5e-10;
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                                                                                             ; TYPE: PRT
; ORGANISM: Human immunodeficiency virus
US-09-576-824A-160
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/146,028
FILING DATE: 22-NOV-1993
INFORMATION FOR SEQ ID NO: 160:
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APPLICATION NUMBER: US/09/680,497
TITING DATE: 06-OCT-2000
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1 Similarity 95.8%;
23; Conservative
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LENGTH: 24 amino acids
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SOFTWARE: Patentin Ver. 2.1
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APPLICANT:
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US-09-680-497-160
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                             SEQ ID NO 160
LENGTH: 24
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Matches 2
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PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
PROCESS FOR THE DETERMINATION OF ANTIBODIES OF
A PROCESS FOR DETERMINATION OF ANTIBODIES OF
BOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
EPITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
CONTAINING THEM
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   Indels
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Sequence 160, Application US/09790497A

GENERAL INFORMATION:

APPLICANT:

TITLE OF INVENTION:

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APPLICANT: De Leys, Robert

TITLE OF INVENTION: PROCESS FOR THE DETERMINANTITLE OF INVENTION: TO INMUNOLOGICALLY IMPORA

TITLE OF INVENTION: A PROCESS FOR DETERMINANTITLE OF INVENTION: BIOITINVLATED PEPTIDES COFTITLE OF INVENTION: CONTAINING THEM

FILE REPERENCE: 2752-11

FURENT APPLICATION NUMBER: US/09/576,824A

CURRENT FILING DATE: 1996-09-30

PRIOR FILING DATE: 1996-09-30

PRIOR FILING DATE: 1993-01-22

PRIOR FILING DATE: 1993-03-08

PRIOR FILING DATE: 1992-03-06
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Human immunodeficiency virus US-09-790-497A-160
                                                               1 NNTRKSERIORGPGRAFVTIGKIG 24
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                                                                                                                           NNTRKSIRIORGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 160, Application US/09576824A; Patent No. 6667387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 95.84
Matches 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Ver. 2.1
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 9
US-09-576-824A-160
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23;
Matches
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us-08-869-386-3.rai

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UTFC305PCT
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                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 512-320-7200
TELEFAX: 512-474-7577
TELEX: NOC Applicable
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
TYPE: AMINO ACID
         32,165
                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 95.8%;
Matches 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 840-333
TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 93.5
Best Local Similarity 95.8
Matches 23; Conservative
                                                                                                                                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: New York
STATE: New York
COUNTRY: USA
ZIP: 10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                            PCT-US92-10378-3
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APPLICANT: BOARD OF REGENTS, THE UNIVERSITY OF
APPLICANT: SASTRY, Jagannadha K.
APPLICANT: ALLNGHAUS, Ralph B.
APPLICANT: ALLNGHAUS, Chris D.
APPLICANT: NEHETE Pramed N.
TITLE OF INVENTION: METHODS AND COMPOSITIONS
TITLE OF INVENTION: FOR ELICITING IMMUNE OR ANTI-INFECTIVE RESPONSES
THILE OF INVENTION: FOR ELICITING IMMUNE OR ANTI-INFECTIVE RESPONSES
ADDRESSEE: Arnold, White & Durkee
STREET: P.O. BOX 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.5%; Score 115; DB 5; Length 24; 95.8%; Pred. No. 2.5e-10;
COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: 13.5" Moiskette, 1.44 Mb
COMPUTER: 18M PS/2 Model 50Z or 55SX
OPERATING SYSTEM: 18M P.C. DOS (Version 3.30)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/06688
FILING DATE: 19920811
CLASSIFICATION NUMBER: 744 281
FILING DATE: 13 ANGUSE 1992
ATTORNEY AGENT INFORMATION:
FILING DATE: 13 ANGUSE 1992
ATTORNEY AGENT INFORMATION:
FILING DATE: 13 ANGUSE 1992
ATTORNEY AGENT INFORMATION:
FELEPANCE (617) 542-8906
TELEPHONE: (617) 542-8906
TELEPHONE: (617) 542-8906
INFORMATION FOR SEQ ID NO: 12:
LENGTH: 24 ANTERISTICS:
LENGTH: 24 ANTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
RIPLICATION NUMBER: PCT/US92/10378
FILING DATE: 19921202
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/800,932
FILING DATE: December 2, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/945865
FILING DATE: September 16, 1992
ATORNEY/AGENT INFORMATION:
NAME: PATKEY, DAVIG L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 NNTRKSIRIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCT-US92-10378-3; Sequence 3, Application PC/TUS9210378; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 95.81
Matches 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: AMINO ACID TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCT-US92-06688-12
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Score 115; DB 5; Length 24; Pred. No. 2.5e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93.5%; Score 115; DB 3; Length 25; 95.8%; Pred. No. 2.6e-10; tive 0; Mismatches 1; Indels
                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER FIGURAL FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: BATCH FOLDS/MS-DOS
SOFTWARE: PATCHIN Rolease #1.0, Version #1.25
SOFTWARE: PATCHIN Rolease #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,324
FILING DATE: 09-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-APR-1997
ATTORNEY AGENT INFORMATION:
NAME: Evans, Barry
REGISTRATION NUMBER: 370132-2000
TELERDHOME: (212) 840-3333
TELERDHOME: (212) 840-3333
                                                 0; Mismatches
                                                                                                                                                                                                                                      US-08-485-324-13
Sequence 13, Application US/08485324
Fatent No. 6043093
GENERAL INFORMATION:
APPLICANT: Wohlstadter, Jacob
INTILE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: Curits, Worris, & Safford
ADDRESSEE: Clo Barry Evans
STREET: 530 Fifth Avenue
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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                      CAPPUTER FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: Eloppy disk
COMPUTER: Elam PC Compatible
OPERATIONS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/447,506
FILING DATE: 23-MAY-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/235,437
RILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTONEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: 23-MAY-1995
CLIANG DATE: 23-MAY-1995
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 08/235,437
FILING DATE: LOS APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: LOS APR-1992
ATTORNEY/AGENT INFORMATION:
ANALY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 115; DB 3;
Pred. No. 2.6e-10;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                        NAME: EVANG, BAITY
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
TELECOMUNICATION INFORMATION:
TELECHONE: (212) 840-333
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 31, Application US/08447506
Patent No. 6066499
GENERAL INFORMATION:
APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: CLUTTIS, MOTTIS, & Safford
ADDRESSEE: C/O Barry Evans
STREET: 230 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 NNTRKSERIORGPGRAFVTIGKIG 24
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REGISTRATION NUMBER: 22,802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 95.8°
Matches 23; Congervative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-447-506-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-447-506-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
93.5%; Score 115; DB 3;
Best Local Similarity 95.8%; Pred. No. 2.6e-10;
Matches 23; Conservative 0; Mismatches 1;
                                                                                                                                                 Sequence 31, Application US/08485324
Patent No. 6043093
GENERAL INFORMATION:
TITLE OF INVENTION: SELECTION METHODS
TITLE OF INVENTION: SELECTION METHODS
CORRESPONDENCE ADDRESS:
ADDRESSEE: CVC Barry Evans
STREET: 530 Fifth Avenue
CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 07-UN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORWATION:
NAME: Evans, Barry
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Curtis, Morris, & Safford
ADDRESSEE: c/o Barry Evans
STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
1 NNTRKSERIQRGPGRAFVTIGKIG 24
                                        NNTRKSIRIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 NNTRKSERIORGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 NNTRKSIRIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13, Application US/08447506
Patent No. 6066499
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (212) 840-3333
TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: linear
US-08-485-324-31
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STATE: New York
                                                                                                                                     US-08-485-324-31
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US-08-447-506-13
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us-08-869-386-3.rai

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NAME: Evans, Barry
REGISTRATION NUMBER: 22,802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 840-712
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 95.89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25 amino acida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                               10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-235-437-31
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                                                                                                                                                                                                                                                                                          93.5%; Score 115; DB 3; Length 25; 95.8%; Pred. No. 2.6e-10; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93.5%; Score 115; DB 3; Length 25; 95.8%; Pred. No. 2.6e-10; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/235,437

FILING DATE: 29-APR-1994

CLASSIFCATION: 435

PROR APPLICATION DATA:

APPLICATION NUMBER: US/0852,412

FILING DATE: 16-ARR-1992

ATTORNEY/AGENT INFORMATION:

NAME: Evans, Barry

REGISTRATION NUMBER: 370132-2000

REFERENCE/DOCKET NUMBER: 370132-2000

TELECOMMUNICATION INFORMATION:

SEQUENCE CHARACTERICATICS:
TELECOMOUNICATION INFORMATION:
TELECOMOUNICATION INFORMATION:
TELEFAX: (212) 840-3333
TELEFAX: (212) 840-712
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris, & Safford
ADDRESSEE: Cortis, Morris, & Safford
ADDRESSEE: Cortis, Morris, Cortis, New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 NNTRKSERIQRGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 10036
COMPUTER READMALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13, Application US/08235437
Patent No. 6087177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 95.8<sup>1</sup>
Matches 23; Conservative
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                                                                                                                                                                                                                                                                                                                                           23; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                             US-08-447-506-31
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Gaps
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                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/235,437
FILING DATE: 29-APR-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION: WHEER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Evans, Barry
NAME: Evans, Barry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patentin Release #1.0, Version #1.25
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95.8%; Pred. No. 2.6e-10;
:ive 0; Mismatches 1;
Sequence 31, Application US/08235437
Patent No. 6087177
GENERAL INFORMATION:
TITLE OF INVENTION: SELECTION METHODS
TITLE OF SEQUENCES: 31
CORRESPONDENCE ADDRESSE:
ADDRESSEE: C/O Barry Evans
STREET: 530 Fifth Avenue
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-447-515-13
; Sequence 13, Application US/08447515
; Patent No. 6162640
; GENERAL INFORMATION:
; APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
; NUMBER OF SEQUENCES: 31
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Curtis, Morris, & Safford
; ADDRESSEE: Curtis, Morris,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 370132-2000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 NNTRKSIRIQRGPGRAFVTIGKIG 24
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
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                                                                                                                                                                           Query Match 93.5%; Score 115; DB 3; Length 25; Best Local Similarity 95.8%; Pred. No. 2.6e-10; Matches 23; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 99, Application US/08257528B
Patent No. 5639854
GENERAL INPORMATION:
APPLICANT: CHONG, Pele
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STRRET: Sulte 701, 330 University Avenue
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 99, Application US/08460602A
Patent No. 5759769
GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/257,528B
FILING DATE: 09-UNN-1994
CLASSIFICATION: 424
ATTORNEY AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-336 MIS: jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                      1 NNTRKSERIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                            1 NNTRKSIRIQRGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopny At
COMPUTED:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24 amino acids
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                                                         single
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                            TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 22; Conserva
                                                                                     linear
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; STRANDEDNESS: sin;
; TOPOLOGY: linear
US-08-257-528B-99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ontario
                                                                                  ; TOPOLOGY:
US-08-447-515-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-257-528B-99
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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MEDIUM TYPE: Floppy disk
COMPUTER: TEM PC compatible
COMPUTER: TEM PC compatible
COMPUTER: TEM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/447,515
FILING DATE: 23-MAY-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 08/235,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: EVADA: BATE: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
TELECOMMONICATION INVERS: 370132-2000
TELECOMMONICATION INVERS: 370132-2000
                                                            FILING DATE: 43-1...
CLASSIFICATION: 435
PRIOR APPLICATION 1947A:
APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY YAGENT INFORMATION:
NAME: Bvans, Barry
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
FILECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-0712
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
INFORMATION COT SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
INFORMATION ACIDS
INFORMATI
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93.5%; Score 115; DB
Best Local Similarity 95.8%; Pred. No. 2.6e
Matches 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 31, Application US/08447515
Patent No. 6162640
GENERAL INPORMATION:
TITLE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: CURTIS, MOXIS, & Safford
ADDRESSEE: CORRESPONDENCE
STREET: S30 Fifth Avenue
CITY: New York
STATE: New York
23-MAX-1995
7N: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 NNTRKSERIQRGPGRAFVTIGKIG 24
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INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-447-515-31
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88.6%; Score 109; DB 1; Length 24;
Best Local Similarity 95.7%; Pred. No. 1.9e-09;
Matches 22; Conservative 0; Mismatches 1; Indels
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US-04-60-21/A-59

US-04-60-21/A-59

Patent No. 5800822

GENERAL INFORMATION:

APPLICANT: CHONG, Pele

APPLICANT: ALDIA Michel H.

TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides

NUMBER OF SEQUENCES: 101

CORRESPONDENCE ADDRESS:

ADDRESSEE: Sim & McBurney

STREET: Suite 701, 330 University Avenue

CITY: Toronto

STATE: Ontario

COMPUTR: Canada

LIP: MGG IR7

COMPUTR: READALE FORM:

MEDIUM TYPE: Floppy disk

COMPUTR: READALE FORM:

MEDIUM TYPE: Floppy disk

COMPUTR: Be Teloppy disk

COMPUTR: Be Teloppy disk

COMPUTR: Be Teloppy disk

COMPUTR: Be Petentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: 08/08/465,217A

FILING DATE: 09-JUN-1994

CLASSIFICATION: 424

PRIOR APPLICATION ATA:

APPLICATION NUMBER: 08/257,528

FILING DATE: 09-JUN-1994

CLASSIFICATION: 424

PRIOR APPLICATION: 424

PRIOR APPLICATION: 424

PRIOR APPLICATION: 424

PRIOR APPLICATION: 424

ATTORNEY/AGATION: 434

                FILING DATE: 05-UN-1993

PRIOR APPLICATION: 424

PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/257,528

FILING DATE: 09-UN-1994

CLASSIFICATION DATA:
APPLICATION NUMBER: 08/073,378

FILING DATE: 09-UN-1993

CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973

REGISTRATION NUMBER: 24,973

RELEPHONE: (416) 595-1163

INFORMATION FOR SEQ ID NO: 99:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids

TYRE: mino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 NTRKSERIORGPGRAFVTIGKIG 24
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     05-JUN-1995
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-463-966A-99
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Pred. No. 1.9e-09;
0; Mismatches 1; Indels
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| Patent No. 5795955
| GENERAL INFORMATION
| PAPLICANT: CHONG, Pele
| APPLICANT: KLEIN, Michel H.
| TILLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
| NUMBER OF SEQUENCES: 101
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Sim & McBurney
| STREET: Sulte 701, 330 University Avenue
                                                             STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/463,966A
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sinf & Meburney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 NTRKSERIQRGPGRAFVTIGKIG 24
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Best Local Similarity 95.7%;
Matches 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 99, Application US/08464329A
| Patent No. 5817754
| GENERAL INFORMATION:
| APPLICANT: GLONG, Pele
| APPLICANT: KLEIN, Michel H.
| TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
| NUMBER OF SEQUENCES: 101
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Sim & McBurney
| STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: IEM PC compatible
COMPUTER: IEM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/464,329A
FILING DATE: 05-JUN-1995
CLASSIPICATION: 424
PRIOR APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/073,378
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
424
PR
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NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-449 MIS:jb
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 NTRKSERIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 NTRKSIRIORGPGRAFVTIGKIG 23
(416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (416) 595-1155
                                   TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                  LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 24 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                    STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                      amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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Matches 22; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: Canada
ZIP: M5G 1R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Toronto
STATE: Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
TELEPHONE:
                                                                                                                                                                                                                                                                                              ; TOPOLOGY:
US-08-465-217A-99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-464-329A-99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-464-329A-99
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                                                                                              APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLBIN, Michel H.
TILLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
TUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:

SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/462,507A
FILING DATE: 0-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 0474
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION INFORMATION:
MET STEWART NUMBER: 1038-451 MIS:jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 99, Application US/08467881A
Patent No. 5951986
GENERAL INFORMATION:
APPLICANT: CHONG, Pele
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                           ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                         CONTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS.
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US-08-462-507A-99
; Sequence 99, Application US/08462507A
; Patent No. 5876731
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 99:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
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US-08-462-507A-99
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Gaps ö

2 NTRKSERIQRGPGRAFVTIGKIG 24

Conservative

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28-JUN-1994
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STATE: Texas
COUNTRY: USA
ZIP: 77025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE:
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APPLICANT: GIRARD, MARC
TITLE OF INVENTION: INDUCTION OF PROTECTION AGAINST VIRAL
TITLE OF INVENTION: INFECTION BY SYNERGY BETWEEN VIRUS ENVELOPE GLYCOPROTEIN
TITLE OF INVENTION: AND PEPTIDES CORRESPONDING TO NEUTRALIZATION EPITOPES OF
TITLE OF INVENTION: THE GLYCOPROTEIN
NUMBER OF SEQUENCES: 23
CORRESPONDENCES: ADDRESS:
ADDRESSEE: FINNERAN, HENDERSON, FARABOW, GARRETT &
ADDRESSEE: DUNNER, L.L.P
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                                                                                                                                                                                                                            CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/467,881A

FILING DATE: 06-JUN-1995

CLASSIFICATION NUMBER: US/08/57,528

FILING DATE: 09-JUN-1994

CLASSIFICATION DATA:

APPLICATION NUMBER: 08/073,378

FILING DATE: 09-JUN-1994

CLASSIFICATION: 424

PRIOR APPLICATION: 424

PRIOR APPLICATION: 424

PRIOR APPLICATION: 424

ATTORNEY AGENT INFORMATION:

NAME: STEWART, MICHAEL I.

REGISTRATION: WUMBER: 24,973

REFERENCE/DOCKET NUMBER: 1038-488 MIS:jb

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.

ZIP: 2005

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/266,448
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 NTRKSERIQRGPGRAFVTIGKIG 24
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                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/08266448
Patent No. 5876724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 1300 I Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPAONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 99:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 24 amino acids
                                                                   STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-467-881A-99
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Fung, Michael S.C., Sun, Bill N.C., Sun, Cecily R.Y.,
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GENERAL INFORMATION:
APPLICANT: Chang, Tae Wen, Fung, Michael S.C., Sun, Bill N.C., Sun, Ceci
APPLICANT: Chang, Nancy T.
TILLS OF INVENTION: Monoclonal Antibodies which Neutralize HIV-1 Infection
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSER: Tanox Biosystems, Inc.
STREET: 10301 Stella Link Rd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Hi Density Diskette
COMPUTER: 1BM PS/2
OPERATING SYSTEM: DOS, Version 3.30
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/950,571A
FILING DATE: 19920922
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: NO. 5854400 07/767,533
FILING DATE: 09/26/1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Meyers, Kenneth J.
REGISTRATION WUMBER: 25,146
REFERENCE/DOCKET NUMBER: 03495.0088-13
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Mirabel, Eric P.
REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TNX87-11BBC
TELECOMUNICATION INFORMATION:
TELEPHONE: 713-664-2288
CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION UNDRER: US 08/145,664

FILING DATE: 04-NOV-1993

PRIOR APPLICATION UNDRER: US 07/782,241

FILING DATE: 28-OCT-1991

PRIOR APPLICATION NUMBER: US 07/672,647

FILING DATE: 18-MAR-1991

PRIOR APPLICATION NUMBER: US 07/494,749

FILING DATE: 19-MAR-1991

PRIOR APPLICATION NUMBER: US 07/494,749

FILING DATE: 19-MAR-1991

APPLICATION NUMBER: US 07/494,749

FILING DATE: 19-MAR-1991

APPLICATION NUMBER: US 07/494,749

FILING DATE: 19-MAR-1991

APPLICATION NUMBER: US 07/494,749

FILING DATE: 19-MAR-1991
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Patent No. 5854400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (202) 408-4132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         not relevant
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Gaps

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Indels

Mismatches

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3 TRKSERIQRGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 236268
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 90.9
Matches 20, Conservative
   21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10016
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                           New York
                                                                                                                                                                                                                                                                                                                                                         New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                         US-08-537-245-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-537-245-1
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   Matches
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                                                                                                                                           Score 105; DB 2; Length 25;
Pred. No. 7.8e-09;
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                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/08345321
Patent No. 5914109
GENERAL INFORMATION:
APPLICANT: ZOLLA-PAZNER, Susan
APPLICANT: GORNY, Miroslav K.
TITLE OF INVENTION: HETEROHYBRIDOMAS PRODUCING HUMAN
TITLE OF INVENTION: MONOCLONAL ANTIBODIES TO HIV-1
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/345,321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 103; DB 2;
Pred. No. 1.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Browdy and Neimark
STREET: 419 Seventh Street, N.W., Suite 300
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZOLLA-PAZNER1B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/872,675
                                                                                                                                                                                                             1 NNTRKSERIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                              4 NNTRKSIRIORGPGRAFVTIGK 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Browdy, Roger L.
REGISTRATION NUMBER: 25,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: ZO
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFRAX: 202-737-3528
                                                                                                                                       85.4%;
illarity 95.5%;
Conservative (
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95.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
INDIVIDUAL ISOLATE: II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 248633
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
 713-664-8914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: 1.22
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide
                                                                   TYPE: AMINO ACID
                                                                                    TOPOLOGY: Linear
                                                                                                                                                        Local Similarity
hes 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                     US-07-950-571A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM:
TELEFAX:
                                                                                                                                                                                                                                                                                                                    US-08-345-321-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-345-321-2
                                                                                                                                         Query Match
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Matches
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Patent No. 603957
Patent No. 603957
APPLICANT: Barl, Patricia L.
APPLICANT: Barder, Christopher C.
APPLICANT: Doms, Robert W.
TITLE OF INVENTION: Oligomeric HIV-1 Envelope Glycoproteins NUMBER OF SEQUENCES: 6
CORRESPONDENCE S.
ADDRESSEE: Knobbe, Martens, Olson and Bear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 101; DB 2; Length 24;
Pred, No. 2.6e-08;
Sequence 1, Application US/08537245
Patent No. 5985275
GENERAL INFORMATION:
APPLICANT: Neurath, A. Robert, Debnath, Asim K.,
APPLICANT: Jiang, Shibo
TITLE OF INVENTION: Proteins and Peptides Modified By
TITLE OF INVENTION: Aromatic Acid Anhydride Compounds
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Frishauf, Holtz, Goodman & Woodward
                                                                                                                                                                                                                                                                                                                                                                                                Diskette, 3+ inch, 0.72 mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Barth, Richard
REGISTRATION NUMBER: 28,180
REFERNCE/DOCKET NUMBER: 950157/RSB
TELECOMMUNICATION:
TELEPHONE: (212) 972-1400
TELEFAX: (212) 370-1622
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/537,245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: cDNA to genomic RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 NNTRKSERIORGPGRAFVTIGK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/420,573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             82.1%;
90.9%;
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us-08-869-386-3.rai

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76.4%; Score 94; DB 2;
    NIH079.1DVCP1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 4, Application US/08452503A; Patent No. 5849475
                                                                                                                                                                                                                                                                                                                                                                                                                       2 NTRKSERIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 NTRKSIRIORGPGRAFVTIGK 22
REPERENCE/DOCKET NUMBER: NIH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-235-8550
TELEFAX: 619-235-0176
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Rovinski, Benjamin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                            LENGTH: 22 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 21 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                       ; FRAGMENT TYPE: internal US-09-070-291-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                     single
                                                                                                                                                                   STRANDEDNESS:
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                                                                                                                                                                                                                                                      ANTI-SENSE:
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US-08-452-503A-4
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Patent No. 6171596
GENERAL INFORMATION:
APPLICANT: Bard, Patricia L.
APPLICANT: Dome, Robert W.
APPLICANT: Dome, Robert W.
APPLICANT: Moss, Bernard
TITLE OF INVENTION: 01igomeric HIV-1 Envelope Glycoproteins NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Knobbe, Martens, Olson and Bear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/805,889
FILING DATE: 03-MAR-1997
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/165,314
FILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/070,291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AUDRESSEE: Knobbe, Martens, Olson and Bear STREET: 620 Newport Center Drive 16th Floor CITY: Newport Beach STATE: CA
      16th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2e-08;
                                                                                                                                                                                                                                                                                                                                                          NAME: Fuller, Michael L.
REGISTRATION NUMBER: 36,516
REFERENCE, DOCKET NUMBER: NIH079.001A
TELECOMUNICATION INFORMATION:
TELEPAX: 619-235-0176
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80.5%; Score 99;
ilarity 95.2%; Pred. No. Conservative 0; Mismatcl
    : 620 Newport Center Drive
Newport Beach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 NTRKSERIQRGPGRAFVTIGK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 36,298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION: NAME: Vensko, Nancy Ways
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE: NO FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION:
                            STATE: CA. 92660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92660
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APPLICANT: Cao, Shi Xian
APPLICANT: Cao, Shi Xian
APPLICANT: Cao, Shi Xian
APPLICANT: Cao, Shi Xian
TITLE OF INVENTION: Retrovirus-Like Particles Containing
TITLE OF INVENTION: Modified Envelope Glycoproteins
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSER: Sim & McBurney
STREET: 330 University Avenue, 6th Floor
CITY: Foronto
Length 22;
                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/452,503A
FILING DATE: J0-MAY-1995
CLASSIFICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION UNBER: US/08/073,526
FILING DATE: 09-JAN-1993
CLASSIFICATION WHERE: US-JAN-1993
CLASSIFICATION WHERE: US-JAN-1993
CLASSIFICATION WHERE: 24,973
REFERENCE/DOCKET NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-447 MIS:a8
TELECOMMULICATION: NIVORMATION:
TELEPHONE: (416) 595-1155
Query Match 80.5%; Score 99; DB 3; 1
Best Local Similarity 95.2%; Pred. No. 5.2e-08;
Matches 20; Conservative 0; Mismatches 1
```

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APPLICANT: ROVINSKI, Benjamin
APPLICANT: ROVINSKI, Benjamin
APPLICANT: PRO, Shi.Xian
APPLICANT: PRO, Shi.Xian
APPLICANT: PRESSON, Roy
APPLICANT: KLEIN, Michel H
TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INPECTIOUS RETROVIRUS-LIKE PARTICLES
CORRESPONDENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSES: Sim & McBurney
SIRRET: 6TH Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                               TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2; Indels
                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC_compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/761,828
                                                                  NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
COMPUTEY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: RIOPPY disk
                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNDAER:
FILING DATE: AUGUST 15, 1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWARY, Michael I
REGISTATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-385 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEFAK: (416) 595-1155
TELEFAK: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 94; DB 2;
Pred. No. 2.7e-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 25, Application US/08761828
Patent No. 5879925
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                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 amino acids
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Best Local Similarity 90.5
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-470-419-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-761-828-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: 1
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  g
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                                 Gaps
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                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Rovinski, Benjamin
APPLICANT: Raynes, Jain
APPLICANT: Cao, Shi Xian
APPLICANT: Cao, Shi Xian
APPLICANT: Klein, Michel H
TITLE OF INVENTION: Retrovirus-Like Particles Containing
TITLE OF INVENTION: Modified Envelope Glycoproteins
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSES: Sim & McBurney
STREET: 330 University Avenue, 6th Floor
COUTY: TOTORIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      76.4%; Score 94; DB 2; Length 21; 90.5%; Pred. No. 2.7e-07; tive 0; Mismatches 2; Indels
                               2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/453,745A
FILING DATE: 30-MAX-1995
CLASSIFICATION DATA:
APPLICATION DATA:
RIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/073,526
FILING DATE: 09-JAN-1993
CLASSIFICATION WINGER: US/08/073,526
FILING DATE: 09-JAN-1993
CLASSIFICATION NUMBER: US/073,526
FILING DATE: 09-SAN-1993
TELECOMUNICATION NUMBER: 24,73
REFERENCE/DOCKET NUMBER: 1038-445 MIS:as
TELECOMUNICATION INFORMATION:
TELEPHONE: (416) 595-1153
INFORMATION FOR EXQ ID NO: 4:
    Pred. No. 2.7e-07;
                          0; Mismatches
                                                                     2 NTRKSERIQRGPGRAFVTIGK 22
                                                                                                                1 NTRKRIRIORGPGRAFVTIGK 21
                                                                                                                                                                                                                              Sequence 4, Application US/08453745A
Patent No. 5866137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 NTRKSERIORGPGRAFVTIGK 22
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Patent No. 586520
GENERAL INFORMATION:
APPLICANT: ROVINSKI, Benjamin
APPLICANT: YAO, Fel-Long
APPLICANT: PERSSON, ROY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 NTRKRIRIORGPGRAFVTIGK 21
  90.5%;
Best Local Similarity 90.5
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 21 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear
US-08-453-745A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ontario
                                                                                                                                                                                   RESULT 35
US-08-453-745A-4
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                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: ROVINSKI, Benjamin
APPLICANT: CAO, Shi-Xian
APPLICANT: YAO, Fai-Long
APPLICANT: YAO, Fai-Long
APPLICANT: PERSSON, ROY
APPLICANT: KLEIN, Michel H
TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
                                                                                Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SIREET: SILCE VOI, 330 ULLVEIBLEY AVELLE
COUNTRY: Canada
ZIP: MGG 1R7
COMPUTER READABLE FORM:
MEDIUM TYRE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,105
FILING DATE: August 15, 1994
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWARY, Michael I
REGISTRATION NUMBER: 24, 973
REFERENCE/DOCKET NUMBER: 1038-385 MIS:jb
TELEFHONE: (416) 595-1153
TELEFRAM: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LEMGTH: 21 maino acids
                                                                            Score 94; DB 2;
Pred. No. 2.7e-07;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 94; DB 2; I
Pred. No. 2.7e-07;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                    Sequence 25, Application US/08290105
Patent No. 5955342
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 25, Application US/08776949;
Patent No. 6025125;
GENERAL INFORMATION:
APPLICANT: Rowinski, Benjamin;
APPLICANT: Cao, Shi-Xian
                                                                                                                                                                    2 NTRKSERIQRGPGRAFVTIGK 22
                                                                                                                                                                                                              1 NTRKRIRIORGPGRAFVTIGK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 NTRKSERIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 NTRKRIRIQRGPGRAFVTIGK 21
                                                                                   76.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            76.4%;
                                                              Query Match
Best Local Similarity 90.5%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
nes 19; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
                      linear
STRANDEDNESS:
                    ; TOPOLOGY:
US-08-452-5208-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 40<sup>(--</sup>
US-08-776-949-25
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                                                                                                                                                                                                                                                                                  RESULT 39
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Sequence No. 5912338

Patent No. 5912338 5840872

GENERAL INFORMATION:

APPLICANT: Rovinski, Benjamin
APPLICANT: Rovinski, Denjamin
APPLICANT: Rovinski, Joel
APPLICANT: Riein, Michel H

TITLE OF INVENTION: Retrovirue-Like Particles Containing
TITLE OF INVENTION: Modified Envelope Glycoproteins
NUMBER OF SEQUENCES: 7

CORRESPONDENCE ADDRESS:
ADDRESSER: Sim & McBurney
STREET: 330 University Avenue, 6th Floor
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 76.4%; Score 94; DB 2; Length 21; Best Local Similarity 90.5%; Pred. No. 2.7e-07; Matches 19; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Ploppy disk
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/452,520B
FILING DATE: 30-MAY-1995
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US/08/073,526
FILING DATE: 09-JAN-1993
CLASSIFICATION: A135
ATTORNEY/AGENT INFORMATION:
NAME: Stewart, Michael I
REGISTRATION NUMBER: 1038-446 MIS:a8
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                     ATTORING DATE: 15-M04-1994
ATTORING DATE: 15-M04-1994
ATTORING DATE: 15-M04-1994
ATTORING THE STEWART, MICHAEL I
NAME: STEWART, MICHAEL I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-655 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1155
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25: SEQUENCE CHARACTERISTICS:
LENGTH: 21 and acide
TYPE: amino acide
TYPE: amino acide
TYPE: amino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 NTRKSERIQRGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 NTRKRIRIORGPGRAFVTIGK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 19; Conservative
                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
                                                                                                                                                                                                                                                                                                                                                                   ;
TOPOLOGY:
US-08-761-828-25
                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Query Match 76.4%; Score 94; DB Best Local Similarity 90.5%; Pred. No. 2.7e Matches 19; Conservative 0; Mismatches
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STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/290,105
FILING DATE: 15-AUG-1994
ATTORNEY AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-798 M
TELECOMMUNICATION INFORMATION:
FILEFAX: (416) 595-1165
FILEFAX: (416) 595-1163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 NTRKSERIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 NTRKRIRIQRGPGRAFVTIGK 21
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Patent No. 6080408

GENERAL INFORMATION:
APPLICANT: ROVINSKI, Benjamin
APPLICANT: ROVINSKI, Benjamin
APPLICANT: PERSSON, ROO, Fei-Long
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES MADE NON-
TITLE OF INVENTION: INFECTIOUS BY A PLURALITY OF MUTATIONS
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & MCBurney
                             APPLICANT: Persson, Roy
APPLICANT: Klein, Michel H
TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: SIM & MCBurney
STREET: 6th Floor, 330 University Avenue
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 76.4%; Score 94; DB 3; Length 21; Best Local Similarity 90.5%; Pred. No. 2.7e-07; Matches 19; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: Canada
ZIP: MSG IR7
COUNTRY: Canada
ZIP: MSG IR7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: TBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version MIL.30
SOFTWARE: Patentin Release #1.0, Version #1.30
STELEST RELEATION NUMBER: 24,973
REGERENOR! DOCKET WUMBER: 24,973
FELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1163
SOFTWARTION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LEMGTH: SOFTWARE
LEMG
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PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 NTRKSERIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 NTRKRIRIORGPGRAFVTIGK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Floppy disk
IBM PC compatible
Yao, Fei-Long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-776-949-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-482-810-25
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Gaps
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| Patent No. 6291157
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| APPLICANT: CAO, Shi.Xian
| APPLICANT: YAO, Fei.Long
| APPLICANT: PERSON, ROY
| APPLICANT: KLEIN Michel H
| TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
| TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
| NUMBER OF SEQUENCES: 26
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Sim & MCBURTEY
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Sim & MCBURTEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Toronto
STATE: Ontario
CONTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Eloppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/027,955
FILLING DATE:
CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 22-AUG-1994
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-490 MIS:vg
TELEPROVIC (416) 595-1163
TELEPROVIC (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 94; DB 3;
Pred. No. 2.7e-07;
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FILING DATE:
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CAO, Shi-Xian
YAO, Fei-Long
PERSSON, ROY
KLEIN, Michel H
TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
RETROVINUS-LIKE PARTICLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: OLICALO
STATE: OLICALO
STATE: OLICALO
COUNTRY: Canada
ZIP: MGG LTF
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: DATE: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,805
FILING DATE: 10-Aug-2000
CLASSIFICATION: CURNOWN>
PRIOR APPLICATION AUTHORNATION:
APPLICATION DATE:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
AND AND AUTHORNATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 76.4%; Score 94; DB 3; Length 21; Best Local Similarity 90.5%; Pred. No. 2.7e-07; Matches 19; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                   Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-1068 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSER: 310 & Meburney
STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                   Query Match 76.4%; Score 94; DB 3; 18est Local Similarity 90.5%; Pred. No. 2.7e-07; Matches 19; Conservative 0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 43
US-09-636-805-25
; Sequence 25, Application US/09636805
; Patent No. 6342228
; GENERAL INFORMATION;
                                                                                                                                                                                                                                                                                                                                       2 NTRKSERIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                            1 NTRKRIRIQRGPGRAFVTIGK 21
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LENGTH: 21 amino acids
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        Bingle
                                                   linear
; STRANDEDNESS:
; TOPOLOGY: lin
US-09-027-955-25
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                    APPLICANT: CAO, Shi-Xian
APPLICANT: YAO, Shi-Xian
APPLICANT: YAO, Fai-Long
APPLICANT: PERSSON, ROY
APPLICANT: KLEIN, Michel H
ATITLE OF INVENTION: INFECTIOUS BY A PLURALITY OF MUTATIONS
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS RETROVIRUS-LIKE PARTICLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 21;
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                                                                                                                                                                                                         ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READARLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/09/258,128
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ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNDRER:
TILNG DATE: 22-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-924 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 25, Application US/09635754
Patent No. 6518030
GENERAL INFORMATION:
APPLICANT: ROVINSKI, Benjamin
YAO, Fei-Long
PRESSON, Fei-Long
KLEIN, Michel H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 NTRKSERIQRGPGRAFVTIGK 22
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 90.5'
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: linear
US-09-258-128-25
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GENERAL INFORMATION:
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DAYA:
APPLICATION NUMBER: US/09/635,754
FILING DATE: 10-Aug-2000
CLASSIFICATION NUMBER: US 09/027,955
FILING DATE: 23.FEBL1998
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 1038-1065 MIS:jb
TELECOMMUNICATION INFORMATION:
TELERRONE (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acid
STYPE: almino acid
STANDEDNESS: single
SEQUENCE DESCRIPTION: SEQ ID NO: 25:
US-09-635-754-25
Mismatches 2;
Mismatches 2;
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Search completed: May 16, 2005, 13:06:19 Job time : 37.9231 Becs

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version 5.1.6	- 2005 -
GenCore	(0) 1993
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OM protein - protein search, using sw model

May 16, 2005, 12:40:16; Search time 169.846 Seconds (without alignments) 54.651 Million cell updates/sec Run on:

US-08-869-386-3 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

2105692 seqs, 386760381 residues Searched:

768190 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match O% Maximum Match 100% Listing first 45 summaries

A\_Geneseq\_16Dec04:\* 1: geneseqp1980s:\* Database

geneseqp1980s:\* geneseqp2000s:\* geneseqp2001s:\* geneseqp2001s:\* geneseqp2002s:\* geneseqp2003bs:\* geneseqp2003bs:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

ore 11155 11155 11155 11155 11155 11155
Score 115 115 115 115 115 115 115 115 115 11
0 1 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1

Aar15058 HIV-1 amp Aar36587 Virus neu Aar36587 Virus neu Aar40475 Human imm Aaw87618 Epitope o Aar42153 gpl20 V3 Aaw07392 HIV-1 str Aar13120 Binding s Aaw72819 HIV-1 gpl Aay68137 HIV-1 III Aaw76842 Fusion im Aav6606 Epitope c Aar3073 Antigent Aav75478 HIV-1 iso Aaw75678 HIV-1 iso Aaw75678 HIV-1 iso Aaw75678 HIV-1 str Aay16052 HIV-1 str Aay16052 HIV-1 str Aay16052 HIV-1 str Aay16052 HIV-1 str Aay57470 HIV BKU V Aac77470 HIV BRU V Aac77470 HIV BRU V	Aar63820 HIV-1 gp1
AAR15058 AAR36587 AAR0475 AAR07153 AAR072153 AAR072153 AAR07219 AAR13120 AAR1219 AAR15137 AAR0566 AAR0566 AAR05669 AAR05669 AAR05699 AAR05699 AAR05699 AAR05699	AAR63820
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## ALIGNMENTS

Protease inhibitor may be use to suppress organ transplant rejection without serious side effects. It may also be used in prevention and therapy of allergy, aplastic anaemia and systemic erythaematodes. See Protease inhibiting peptide immuno-suppressant - used to suppress rejection reaction in organs transplantation. Organ transplant; autoimmune disease; allergy; aplastic anaemia; systemic erythaematodes. Immunosuppressant protease inhibitor. AAR06211 standard; peptide; 24 AA Claim 1; Page 181; 6pp; Japanese. 88JP-00310635. 88JP-00310635. (first entry) (NITL ) NITTO DENKO CORP. WPI; 1990-233739/31. Sequence 24 AA; also AAR06212 JP02157229-A. 07-DEC-1988; 07-DEC-1988; 10-DEC-1990 18-JUN-1990. Synthetic. AAR06211; RESULT 1 

Length 24; Query Match
93.5%; Score 115; DB 2; Lèngth 24
Best Local Similarity 95.8%; Pred. No. 2.8e-09;
Matches 23; Conservative 0; Mismatches 1; Indels 1 NNTRKSIRIQRGPGRAFVTIGKIG 24 1 NNTRKSERIORGPGRAFVTIGKIG 24

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V3 loop; gp41; envelope protein; MN; prototype; virus; varient; HIV; homology; heteroconjugate; enzyme; epitope mapping; replication; conjugate; immunogenic carrier; keyhole limpet hemocyanin; KLH; ovalbumin; succiny] maleimidomethyl.cyclohexanylcarboxylate; SMCC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "Not in the natural sequence of this isolate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heteroconjugate antibody immunogen RP135 (IIIB).
                                                          (SUPE-) INST SUPERIORE DI SANITA.
(CNDR ) CONSIGLIO NAZ DELLE RICERCHE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 NNTRKSERIORGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR29233 standard; peptide; 24 AA
                                                                                                                                                                                                                                      Claim 1; Page 17; 31pp; English.
 92WO-EP000187
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(first entry)
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Best Local Similarity 95.8
Matches 23; Conservative
                                                                                                     Pasti M,
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                                                                                                                                              WPI; 1992-299983/36
                                                                                                   De Rossi A, Past:
Chieco-Bianchi L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 AA;
 29-JAN-1992;
                              30-JAN-1991;
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14-APR-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-HIV sulphated polysaccharide (SPS) can prevent syncytium formation in HIV infected C4 cells. SPS may be isolated by affinity chromatography with the given resin bound peptide fragment RP135. (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prepn. of anti-HIV sulphated polysaccharide - by affinity chromatography using a resin-bound peptide corresp. to a HIV gp. 120 fragment.
                                                                                                                               protein used in isolation of sulphated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence of peptide DB1 determined from the V3 principal neutralising domain (PND) region of HIV-1 strain HTLV-III B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 115; DB 2;
Pred. No. 2.8e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                              Residues 301-324 of HIV gp 120 protein use polysaccharide by affinity chromatography
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page ?; 34pp; English
                             AAR07018 standard; peptide; 24
                                                                                                                                                                                                      Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                                                                                                                                                   (RICH ) MERRELL DOW PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93.5%;
                                                                                                                                                                            HIV; AIDS; ARC; gp120; RP135.
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                                                                                                                                                                                                                                                                                                                         89US-00295856.
                                                                                                   (first entry)
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 95.8 tes 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 Cardin AD, Jackson RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1990-290631/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24 AA;
                                                                                                                                                                                                                                                                                             05-JAN-1990;
                                                                                                                                                                                                                                                                                                                         11-JAN-1989;
05-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-OCT-2003
25-MAR-2003
28-JAN-1993
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                                                                                    24-OCT-2003
18-JAN-1991
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                                                        AAR07018;
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1D AAR2

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DT 24-C

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XX X

XX X
RESULT 2
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The principal neutralizing domain (PND) of HIV-1 corresp. to a 24- amino acid sequence arranged in a loop determined by a disulfide bridge in the third hypervariable region, V3. of the protein gp 120. The central portion of the V3-PND contains a sequence which is highly conserved in different HIV-1 isolated strains, whereas the amino acids flanking this sequence are variable. The antigenic properties of V3 region are known to be virus-specific; antibodies elicited by WN-derived peptide do not neutralize HTLV-III B virus and vice-versa. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to standardise OS field)
Di Bello C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                Synthetic peptide(s) which enhance infectivity of HIV-1 in cellular cultures - are used for determining HIV-1 virus in blood and other biological materials.
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Dettin M,
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Mammano F, Panozzo M,
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us-08-869-386-3.rag

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Similarity
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           Sequence 24 AA;
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04-JUL-1993
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                                                                                                                                                                                                                                                                                                             Synthetic.
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                                Query Match
Best Local (
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                                                                                                               The sequences given in AAR29226-35 represent peptides which were used as immunogens for the production of antibodies against HIV. These peptides may be either unconjugated or conjugated to an immunogenic carrier, eg. a keyhole limpet hemocyanin (KLH) or ovalbumin, using succinyl maleimidomethyl cyclohexanylcarboxylate (SMCC) as a conjugating agent. Viruses containing these or similar sequences may be recognised by the heteroconjugate enzymes of the invention. The antibodies raised against these sequences may be identified by standard epitope mapping techniques. These antibodies are capable, even at low concentrations, of nearly eliminating viral replication of different strains of HIV. (Updated on 25
                                          Hetero-conjugate antibodies for treating HIV infections - comprise an antibody specific for an effector cell surface antigen and an antibody to a V3 loop of GP-120 envelope protein of HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequence is that of peptide IIIB, derived from the V3 region of HIV gp120, it was used in binding assays for anti-gp120 antibodies. The anti-gp120 antibodies are useful in vaccine formulations for the treatment or prevention of HIV infection. See also AARAS867-R26873. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anti-idiotype antibodies and methods for their selection - useful as vaccines for the prevention and treatment of HIV infection.
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                                                                                                                                                                                                                                                                                    93.5%; Score 115; DB 2; Length 24; 95.8%; Pred. No. 2.8e-09; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus; AIDS; anti-gp120 antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV gp120 V3 region binding assay peptide IIIB.
                                                                                                                                                                                                                                                                                                                                                          24
                                                                                                                                                                                                                                                                                                                                    1 NNTRKSERIQRGPGRAFVTIGKIG 24
                                                                                            Disclosure; Page 19; 69pp; English
                                                                                                                                                                                                                                                                                                                                                         NNTRKSIRIORGPGRAFVTIGKIG
                                                                                                                                                                                                                                                                                                                                                                                                                     AAR26870 standard; peptide; 24 AA
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                                                                                                                                                                                                                                       -MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 95.8°
Matches 23, Conservative
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Potts BJ;
                       WPI; 1992-415475/50.
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                                                                                                                                                                                                                                                               Sequence 24 AA;
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06-MAR-1992;
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20-MAY-1998
Higgins PJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence of peptide B1 which comprises AAs 308-331 from the V3 region of HIV-1 isolate IIIb.
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                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nine peptides from the V3 regions of HIV-1 isolates IIIb, RF and MN
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   Length 24;
                                                             1; Indels
Score 115; DB 2;
Pred. No. 2.8e-09;
0; Mismatches 1;
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                                                                                                                                                            1 NNTRKSIRIQRGPGRAFVTIGKIG 24
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                                                                                                                       1 NNTRKSERIORGPGRAFVTIGKIG
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      93.5%;
95.8%;
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(first entry)
                                                             23; Conservative
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nes 23; Conserv
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29-JAN-1998;
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                                                                     05-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                    HIV gp120 V3 loop-derived peptides (AAR38170-87) are successful in generating CTL responses, esp. peptide R15K (AAR38187); the T-helper cell-inducing peptide includes the sequence C194 (AAR38164); HIV infection-inhibiting peptides are given in AAR38188-206, and are esp. peptides R15K, N24G, E13V, R8K, T13Q and H13N (AAR38165-69). The peptides may also be derived from an influenza virus protein or a sendai virus protein (AAR41014-15). It was observed that peptide N24G (amino acids 308-311), with sequences derived from the V3 loop of HIV-1 IIIB, inhibits HIV-1 infection of primary human T cells by 92% at 1 microg/ml (ca. 0.4-0.6 microw). (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG -2003 to correct D8 field.)
                                                                                                                                                                                                                                                                                                          Peptide composition for treating and preventing viral infections - comprise CTL-inducing epitope and HIV infection-inhibiting sequence or T helper cell-inducing sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV; LDL; low density lipoprotein; human; immune response; infection; immunodeficiency; neoplastic tissue; myalgic encephalomyelitis; ME; viral infection fatigue syndrome; tuberculosis; hepatitis; AIDS; ARC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                             gp120; HIV-1; cytotoxic T-lymphocyte; CTL; T-helper; AIDS; infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93.5%; Score 115; DB 2; Length 24; 95.8%; Pred. No. 2.8e-09; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                    Sastry JK, Arlinghaus RB, Platsoucas CD, Nehete PN;
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                                                                                                                                                                                                                                                                                                                                                     Claim 19; Page 95; 130pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY22581 standard; peptide; 24 AA
           AAR38165 standard; peptide; 24 AA
                                                                                                                                    Human immunodeficiency virus 1.
                                                                                                                                                                                               92WO-US010378
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92US-00945865
                                                   (revised)
(revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23; Conservative
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                                                                                            V3 loop peptide N24G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24 AA;
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16-SEP-1992;
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19-0CT-1999
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                                                    27-AUG-2003
25-MAR-2003
                                                                      12-OCT-1993
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                               AAR38165;
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 RESULT 8
AAY22581
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This sequence represents a HIV sequence that binds human low density conjugated (LDL), and is designated sequence "A". The invention relates lipoprotein (LDL), and is designated sequence "A". The invention relates to a method for enhancing the immune response in a parient with a condition, selected from immunodeficiency (due to a viral, bacterial, wycoplasmic, fungal or parasitic infection, or from the growth of composition farigue syndrome, tuberculosis, or hepatitis. The method comprises using a pharmaceutical composition, comprising a recombinant composition, comprising a recombinant composition, compositions for: the treatment of acquired immunodeficiancy syndrome compositions for: the treatment of acquired immunodeficiancy syndrome (AIDS) or ARC (AIDS related complex); reducing syncytium formation in HIV infected CD4 cells; treating blood or body fluid or organs to contract the human or animal host. The human recombinant LDL receptor in a human or animal host. The human recombinant LDL receptor is a lightly specific inhibitor of HIV-1 replication in vitro. (Updated on 17-OCT-2003) to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Enhancing the immune response using a recombinant human low-density lipoprotein receptor, useful for treating viral infections, especially human immunodeficiency virus (HIV) infection.
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acquired immunodeficiency syndrome; AIDS related complex; HIV-infected CD4 cell; immunosuppressive peptide.
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95.8%; Pred. No. 2.8e-09;
ive 0; Mismatches 1;
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                                                                                                                   Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PREN/) PRENDERGAST P
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The present invention relates to a liposome composition comprising liposomes having an outer surface layer of polyethylene glycol chains, each having a free distal end. A polypeptide or polyaccharide effector molecule is covalently attached to a portion of the distal ends. The effector interferes with specific binding of pathogen or cell in a cffector interfere with specific binding of pathogen or cell in a consolition may be used in treating a condition mediated by binding a pathogen or cell in the bloodstream, to a target cell or cell marrix, and is rapidly removed by cincarding a pathogen or cell in the bloodstream, to a target cell or cell matrix. It can be used in treating septic shock, toxic shock, colonic inflammation, leukaemic cell proliferation, or HIV infection. The present sequence is a peptide of the V3 loop of HIV envelope protein gpl20. This peptide may be used in the composition of the present invention. gpl20 binds to the CD4 receptor during HIV infection of lymphocytes. By cincroducing the present peptide, the CD4 receptors are blocked, thereby childly HIV infection. (Updated on 11-SEP-2003 to standardise OS
                                                                                          Liposome composition for use in treating septic shock comprises liposomes having an outer surface layer of polyethylene glycol chains, and a polypeptide or polysaccharide effector molecule.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93.5%; Score 115; DB 4; Length 24; 95.8%; Pred. No. 2.8e-09; ive 0; Mismatches 1; Indels
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                  Barenholz
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                Martin FJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Simian-Human immunodeficiency virus.
Unidentified.
                                                                                                                                                                            Disclosure; Fig 13; 32pp; English.
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                  Woodle MC,
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Petro J, Okeeffe T;
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                                                        WPI; 2001-201897/20
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                  Zalipsky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                            New peptide 3, amide and heterocyclic compounds and saccharide conjugates used for inhibiting chemokine induced activity and for treating e.g. stroke, vascular diseases, autoimmune diseases and tumor growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    autoimmune diseases, atherosclerosis, osteoporosis, HIV infection and AIDS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotoxaemia, myocardial infarction, acute isotaemia and rheumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in gene therapy for the same diseases, as well as in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chemokines which can be used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences AAB15785-B15794, AAB15803-B15813 and AAB15831-B15848. These chemokine derivatives can be used to treat diseases such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention concerns the identification of a number of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.5%; Score 115; DB 3; Length 24; 95.8%; Pred. No. 2.8e-09; ative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gp120 V3 loop; liposome composition; HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 NNTRKSERIQRGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure, Fig 18; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB68602 standard; peptide; 24 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus 1
                                                                                                                                                         99US-00229071.
99US-00271192.
99US-00452406.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV gp120 V3 loop peptide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93US-00035443.
                                                                                                                   12-JAN-2000; 2000WO-US000821.
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                                                                                                                                                                                                                                                                                Tatalick LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 95.8
Matches 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SEQU-) SEQUUS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
                                                                                                                                                                                                                                                                                                                        WPI; 2000-499101/44.
                                                                                                                                                                                                                                        (NEOR-) NEORX CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 AA;
                                     WO200042071-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                animal models
                                                                                                                                                                                                                                                                                Grainger DJ,
                                                                                                                                                         12-JAN-1999;
17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-MAR-1993;
29-SEP-1994;
                                                                                                                                                                                                   01-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1995;
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25-APR-2001
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                                                                             20-JUL-2000
Synthetic.
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RESULT 10

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HIV 

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Gaps

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Lynn D;

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1 NNTRKSIRIQRGPGRAFVTIGKIG 1 NNTRKSERIQRGPGRAFVTIGKIG

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AAR31276 standard; peptide; 25

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RESULT 13
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                                                           Protein derivative stimulates a lymphocyte proliferative response in HIV-infected humans, providing a means of diagnosis, protection and therapeutic value. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PA field.) (Updated standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conjugate immunogen for AIDS and ARC treatment - composed of neutralising determinant of HIV and Neisseria outer membrane.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This peptide is derived from the HIV IIIB isolate and is cross- reactive with the HIV major neutralisation determinant (MNtD). This MNtD is used in a conjugate, covalently linked to the outer membrane protein (Gmp) from Neisseria, as an immunogen for vaccination against AIDS. A cocktail of different MNtD poly- peptides can be used. See also AAR08274-75 and
                                                                                                                                                                                                                                       Gaps
New HIV proteins and peptide(s) - used in diagnosis, prophylaxis or therapy of AIDS, esp. for prepn. of vaccines against HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunogen; Neisseria outer membrane protein;
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                                                                                                                                                    Revised record issued on 09-SEP-2004 : Correction to location
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 115, DB 2; Length 25;
Pred. No. 3e-09;
0; Mismatches 1; Indels
                                                                                                                                                                                                          Score 115; DB 1; Length 25;
Pred. No. 3e-09;
                                                                                                                                                                                                                                       1; Indels
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                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                 24
                                                                                                                                                                                                                                                                               1 NNTRKSIRIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV major neutralisation determinant
                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV peptide fragment (IIIB isolate).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Emini EA, Marburg S, Scolnick EM,
                                                                                                                                                                                                                                                                 1 NNTRKSERIORGPGRAFVTIGKIG
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                                      Claim 1; Page 27; 29pp; English
                                                                                                                                                                                                                                                                                                                                                            AAR08276 standard; protein; 25
                                                                                                                                                                                                          93.5%;
95.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus.
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89US-00362178,
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ilarity 95.8%;
Conservative (
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                                                                                                                                                                                                                                    23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AIDS; ARC; conjugate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1990-370100/50.
                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                              Sequence 25 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MERI ) MERCK
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06-JUN-1989;
06-JUN-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JUN-1989;
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Best Local S
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/note= "bonds to the OMPC of the conjugate via this site"

Location/Qualifiers

Modified-site

Synthetic

AIDS; ARC; human immunodeficiency virus; vaccine; Neisseria; meningitidis b; outer membrane protein complex; OMPC; PND135.

HIV principal determinant peptide

(first entry)

12-FEB-1993

AAR31276;

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The invention relates to a co-conjugate comprising an immunogenic protein or protein complex having a first set of covalent linkages to low molecular weight moieties which have an anionic or polyamionic character at physiological ph, and a second set of covalent linkages to peptides comprising HIV principal neutralizing determinants (PND's) or immunologically equivalent peptides. Preferably at least one set of the covalent linkages is comprised of maleimide derivatives; the (poly)anionic moiety is composed of one to five residues of the anionic form of a carboxylic, sulphonic or phosphonic acid; the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b, and the PND peptide has a linear structure, a disulphidebonded cyclic structure, an amide-bonded cyclic structure or a thioether-bonded cyclic structure. The present sequence (PND135) is an example of a proper protein companies of the present sequence (PND135) is an example of a proper protein companies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PND peptide component used in the co-conjugate. The co-conjugate is useful for inducing anti-peptide immune response in mammals, for inducing HIV-neutralizing antibodies in mammals, for formulating vaccines to prevent HIV infection or disease, including AIDS, or for treating humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps

    useful for

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conjugate proteins comprising HIV peptide components - usefu preparing vaccines for e.g. AIDS or for treating infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93.5%; Score 115; DB 2; Length 25; 95.8%; Pred. No. 3e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Emini EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            afflicted with HIV infection or disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 NNTRKSERIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 56; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                90US-00555339.
90US-00555966.
91US-00715276.
91US-00715278.
                                                                                                                                                                                                                                                                                                                               91EP-00306598
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Leanza WJ, Marburg S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1992-026505/04
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                                                                                                                                                                                                                                                                                                                             19-JUL-1991;
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                                                                                                                                                                                                                                                            EP467700-A.
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Gaps

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Similarity

23;

Matches

us-08-869-386-3.rag

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The invention relates to a conjugate of an HIV principal neutralizing determinant (PND), or an immunologically equivalent peptide (PEP), covalently coupled to an immunogenic protein or protein complex through an anionic polysaccharide linker. Pref. the immunogenic protein is the outer membrane protein complex (GMPC) of Neisseria meningitidis b and the PND peptide has a linear structure, a disulphide-bonded cyclic structure, an amide-bonded cyclic structure. The present sequence (PNDI35) is an example of a PND peptide component. The conjugates are used for inducing HIV-neutralising antibodies or for making vaccines to prevent contraction of HIV infection or disease. The antibodies can be used for passively protecting against infection by HIV, or for protecting against proliferation of HIV post-infection, or for treating AIDS, or in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV peptide-polysaccharide-protein conjugates - used in vaccines or to produce antibodies to prevent or treat HIV infection.

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    /note= "Joins onto polysaccharide-protein complex via

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV-1; human immunodeficiency virus; antibody generation; AIDS;
infection; CD4 binding site; soluble CD4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93.5%; Score 115; DB 2; Length 25; 95.8%; Pred. No. 3e-09; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV gp120 V3 loop immunogenic peptide RP135 (IIIB).
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                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR33222 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                  Emini EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 9; Page 57; 63pp; English.
                                                                                                                                                                                                                                                      90US-00555974.
91US-00715275.
91US-00715277.
                                                                                                                                                                                                 90US-0055558
                                                                                                                                                                                                                                       90US-00555558.
                                                                                       this site"
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 95.8
Matches 23, Conservative
                                                                                                                                                                                                                                                                                                                                                                  Marburg S, Tolman RL,
                                                                                                                                                                                                                                                                                                                              (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1992-034437/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 25 AA;
                                       Key
Modified-site
                                                                                                                                                                                                     19-JUL-1990;
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                                                                                                                             EP468714-A.
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 Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used as part of a conjugate comprising the major immune enhancing protein (MIEP) of Neisseria meningitidis covalently linked to the HIV PND. The conjugate may be used to prepare vaccines against HIV infections, e.g. AIDS, as research tools for studying PND structure- function relationships, or as immunogens for use in the passive immunisation of humans. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                           Human immunodeficiency virus; AIDS; PND; MIEP; conjugate; major immune enhancing protein; vaccine; anti-HIV antibodies; immunogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide is HIV principle neutralising determinant (PND) 135 and is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conjugates of HIV-1 PND peptide(s) with the MIEP of Neisseria meningitidis - useful as a vaccine for treating and preventing HIV-1 infection, e.g. AIDS in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus; principal neutralizing determinant; outer membrane protein complex; OMPC; Neisseria; AIDS; PND135.
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Pred. No. 3e-09;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                           HIV principle neutralising determinant 135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tolman RL;
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95.8%; Pred
0; }
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                                                                                                            AAR30031 standard; peptide; 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Emini A, Liu MA, Marburg S,
                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                             92EP-00201693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               91US-00715273
                                                                                                                                                                                      (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                               passive immunisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-425771/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-JUN-1991;
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                                                                                                                                                                                    25-MAR-2003
28-APR-1993
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                                                                                                                                                  AAR30031;
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Matches

RESULT 15

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us-08-869-386-3.rag

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The HIV gp120 epitope is used to isolate, create or evolve novel mols. including (in) organic and biomolecules such as proteins, peptides, nuclaids acids, oligonucleotides, lipids, and polyaeaccharides for use as reactants, catalysts, enzymatic cofactors, repressors, enhancers, hormones and binders for a wide variety of substrates in industrial and therapeutic products. This epitope was isolated from variable region 3 of HIV gp120 (amino acids 271-295). (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                The sequences given in AAR41336-39 and AAR42664 represent peptides derived from the V3 region of HIV envelope gp120. These peptides may be used in a vaccine which is effective in humans and animals and activates humoral and cellular immunity. The vaccine also contains a carrier protein containing a cysteine group, eg. human serum albumin (HSA), keyhole limpet haemocyanin (KHM) or multiple antigen peptide. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 24-OCT-2003 to standardise
                                  Virus for prevention of HIV infected diseases – comprising several peptide(s) consisting of V3 region peptide of envelope Gp., 120, etc. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Obtaining a novel mol. - capable of a desired interaction with a substrate of interest and a selection molecule expressed by the host.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV; haemagglutinin; reactants; catalysts; cofactors; repressors; enhancers; hormones; binders; human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                   2; Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                                 Score 115; DB 2;
Pred. No. 3e-09;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 NNTRKSIRIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 151; Page 147; 165pp; English.
                                                                                                              Disclosure; Page 3; 35pp; Japanese
                                                                        complex including carrier protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                           1 NNTRKSERIORGPGRAFVTIGKIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR41330 standard; peptide; 25
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                                                                                                                                                                                                                                                                                                                                                                   93.5%;
95.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus.
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                         23; Conservative
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WPI; 1993-320455/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1993-320737/40.
                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV gp120 epitope
                                                                                                                                                                                                                                                                                                                               Sequence 25 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-MAR-1993;
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21-APR-1994
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                                                                                                                                                                                                                                                                                       OS field)
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence is that of peptide RP135 (IIIB) used as an immunogen for the generation of antibodies directed against the V3 loop of HIV gp120. These antibodies can be used as part of a compsn. With antibodies directed against the CD4 binding site of gp120. The antibodies act synergistically to neutralise HIV-1 in the treatment of HIV infection caused by different strains. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                       Synergistic compsn. for treating HIV-1 infection - comprises antibody to V3 loop of GP120 and antibody to CD4 binding site of GP120 or soluble CD4 polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               V3 region; HIV; envelope gp120; vaccine; human; humoral response; cellular immunity; carrier protein; human serum albumin; HSA; keyhole limpet haemocyanin; KLH; multiple antigen peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93.5%; Score 115; DB 2; Length 25; 95.8%; Pred. No. 3e-09; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                Herlihy WC,
                                                                                                                                                                                                                                Potts BJ, Whitescharf ME, Field KG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV gp120 V3 region peptide HIV-III B.
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92JP-00237648.
93JP-00054239.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sest Local Similarity
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14-AUG-1992;
15-MAR-1993;
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                                                                                                                                  09-SEP-1991;
                  WO9304693-A1
                                                                                            02-SEP-1992;
                                                                                                                                                      20-JUL-1992;
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                                                        18-MAR-1993
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RESULT 17 AAR41336

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                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV-IIIB; peptide 135; principal neutralising domain; antibodies; diagnosis; prophylaxis; therapy; AIDS.
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Score 115; DB 2; Length 25; Pred. No. 3e-09;
                                                                                  1; Indels
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                                                                                  0; Mismatches
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                                                                                                                                                                    1 NNTRKSERIORGPGRAFVTIGKIG 24
                                                                                                                                                                                                                   1 NNTRKSIRIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR04427 standard; peptide; 25 AA
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89US-00359543.
89US-00407663.
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1 Similarity 95.8%;
23; Conservative
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Lynn DU, Petrobre J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-1989;
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20-SEP-1990
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    Query Match
Best Local (
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                                                                                  Matches
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MARO427

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This sequence represents a variant of the HIV sequence that binds human low density lipoprotein (LDL), and is designated sequence "A" (see AY22581). The sequence "A" peptide is isolated from HIV isolate IIIB(BHB).

IIIB(BHID), and this sequence was isolated from HIV isolate IIIB(BHB).

The invention relates to a method for enhancing the immune response in a patient with a condition, selected from immunodeficiency (due to a viral, bacterial, mycoplasmic, fungal or parasitic infection, or from the growth of neoplastic tissue), myalgic encephalomyelitis (ME), post inoculation of neoplastic tissue), myalgic encephalomyelitis (ME), post inoculation of viral infection fatigue syndrome, tuberculosis, or hepatitis. The method comprises using a pharmaceutical composition, comprising a recombinant LDL receptor or a mimic molecule to the cysteine rich domain of LDL receptor or a mimic molecule to the cysteine rich domain of LDL receptor forms pharmaceutical compositions for: the treatment of acquired immunodeficiency syndrome (AIDS) or RRC (AIDS related complex); reducing syncytium formation in HIV-infected CD4 cells; treating blood or body fluid or organs to neutralise-remove immunosuppressive peptides and/or viruses; or treating hapatitis A, B or C. The pharmaceutical composition also treat a viral infection in a human or animal host. The human cerombinant LDL receptor is also useful for manufacturing medicaments for receptor is a highly specific inhibitor of HIV-1 replication in vitro.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enhancing the immune response using a recombinant human low-density lipoprotein receptor, useful for treating viral infections, especially human immunodeficiency virus (HIV) infection.
                                                                                                                                                                                                                                                                                                                               HIV; LDL; low density lipoprotein; human; immune response; infection; immunodeficiency; neoplastic tissue; myalgic encephalomyelitis; ME; viral infection fatigue syndrome; tuberculosis; hepatitis; AIDS; ARC; acquired immunodeficiency syndrome; AIDS related complex; HIV-infected CD4 cell; immunosuppressive peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                          HIV LDL binding peptide, sequence "A" variant.
1 NNTRKSIRIQRGPGRAFVTLGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 12; 24pp; English.
                                                                                                                                AAY22583 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-IB000149.
                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PREN/) PRENDERGAST P T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-494040/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prendergast PT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9938524-A2
                                                                                                                                                                                                                     17-0CT-2003
19-0CT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-AUG-1999
                                                                                                                                                                              AAY22583;
                                                                                                              AAY22583
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Gape

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Score 113; DB 2; Length 25; Pred. No. 5.7e-09; 1; Mismatches 1; Indels

91.9%;

Query Match Best Local Similarity 91.74 Matches 22; Conservative

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1 NNTRKSERIQRGPGRAFVTIGKIG 24

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Either the N-terminal (a) or C-terminal (b), but not both, may be omitted; either (a) or (b) may comprise any of the following: cysteine, a protein or other moiety capable of enhancing immunogenecity, a peptide from an HIV principal neutralising domain, peptide capable of stimulating T-cells, or general immune stimular. See also ARR04427-R04506 and AAQ04273-Q04279. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                    HIV; therapy; AIDS; principal neutralising domain; antibodies; diagnosis; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Revised record issued on 09-SEP-2004 : Correction to Feature Table Key
                                                                                                                   Cpd. eliciting, binding with neutralising antibodies to HIV variants.
                                                                                                                                                                                                                                                                                                                                                                                                                      Principal neutralising domain of HIV variants - used for producing peptide(s) and antibodies for diagnosis, prophylaxis and/or therapy therapy of HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2; Length 23
                                                                                                                                                                                                                                                                                                                                                              Farley J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88.6%; Score 109; DB 2, 95.7%; Pred. No. 2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                             Javaherian K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 27 (d); Page 84; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR33190 standard; peptide; 24 AA
                        Z
                        AAR04502 standard, protein; 23
                                                                                                                                                                                                                                                                                    88US-00352949.
89US-00359543.
89US-00407663.
                                                                                                                                                                                                                                                              88US-00252949.
                                                                       (revised)
(revised)
(first entry)
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Best Local Similarity 95...
Best Local Similarity 95...
                                                                                                                                                                                                                                                                                                                                                             , Putney SD,
Petrobre J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
                                                                                                                                                                                                                                                                                                                                     (REPK ) REPLIGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1990-147824/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 23 AA;
                                                                                                                                                                                                                                                                                    03-OCT-1988;
01-JUN-1989;
19-SEP-1989;
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                                                                                                                                                                                                                                                              03-OCT-1988;
                                                                                                                                                                                                             WO9003984-A.
                                                                                                20-SEP-1990
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                                                                        09-SEP-2004
25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                               Rusche JR,
                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                           Lynn DU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR33190;
                                                 AAR04502;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 22
AAR33190
ID AAR33
XX
AC AAR33
XX
DT 25-MA
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Grimaila R;

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Gapa

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1; Indels

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Two peptides, gp41[8] and V3 loop derived from the gp120 region of HIV-1 were synthetised and purified. To improve the solubility of the gp41[8] peptide the sequence RSS was added to the amino terminal to produce peptide R-S-Sgp41[8]. The immunodominant HIV-derived peptides were used to ascertain whether E.coli outer membrance protein TraT augments the in vitro T-cell proliferative responses. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunogen; vaccine; HIV-1; T-cell; B-cell; epitope; core protein; gpl20;
                                                                                                                                                                                                                                                                        Compsns. contg. E.coli outer membrane proteins TraT, OmpA or OmpF -
increase immune response and are used for treating auto:immune diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                            Cooper DA
                                              AIDS; HIV; therapy; autoimmune disease; gp120; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 109; DB 2;
Pred. No. 2.1e-08;
0; Mismatches 1.
                      Sequence of HIV-1 derived V3 loop peptide
                                                                                                                                                                                        (BIOT-) BIOTECH AUSTRALIA PTY LTD.
(SVIN-) ST VINCENT'S HOSPITAL SYDNEY LTD.
                                                                                                                                                                                                                           Bell SJD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 NTRKSERIORGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW67414 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                         Example; Page 13; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus 1.
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94US-00257528.
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                                                                                                                                                                                                                            Geczy AF, Russell-Jones GJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.6%;
                                                                                                                                            92WO-AU000423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 95.75
Worker 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1 peptide epitope BRU
11-JUL-1993 (first entry)
                                                                                                                                                                                                                                                  WPI; 1993-093727/11.
                                                                                                                                                                                                                                                                                                  AIDS, cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 AA;
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                                                                                                                                            10-AUG-1992;
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                                                                                             409303762-A1
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                                                                                                                    14-MAR-1993.
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                                                                                                                                                                                                                                                                                                                                                                                                                      PN field.)
                                                                       synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW67414;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              V3 100p
                                                                                                                                                                                                     (SVIN-)
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06-JUN-1995;
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                                                                                                                                                                                                                           17-OCT-2003
                                                                                                                                                                                                                                     26-NOV-1999
                                                                                                                                                                                                                                                                                                                                     US5951986-A.
                                                                                                                                                                                                                                                                                                                                                         14-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Klein MH,
                      invention
                                                                                                                                                                                                       AAY39769;
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                                                              Query Match
                                                                         Best Local
                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a synthetic peptide comprising an amino acid sequence containing a T-cell epitope of an HIV gag protein linked a its C terminus to an amino acid sequence containing a B-cell epitope of an HIV gp41 protein and containing the amino acid sequence: XILKDWX2; where XI = E, A, G or Q, and X2 = A or T, or an amino acid sequence capable of eliciting an HIV-specific antiserum and recognizing the sequence XILKDWX2. The synthetic peptide is useful in vaccines against
                                                              - containing T-cell
                                                                                                              The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-cell epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitopes are generally designed based on the p24 core protein and the B-cell epitopes from the V3 loop of the gpl20 protein from various HIV-1 strains. This sequence corresponds to an HIV-1 B-cell peptide epitope used to immunise a guinea pig
                                                                                                                                                                                                                                                 Gaps
                                                              Synthetic human immunodeficiency virus-1 peptide(s) - containing T-epitope and B-cell epitope(s) are candidate vaccines against HIV-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic chimeric HIV polypeptides - comprising gag protein T-cell epitope linked to gp41 B-cell epitope.
                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-1; human immunodeficiency virus; vaccine; T-cell epitope;
gag protein; B-cell epitope; gp41 protein; chimeric; infection.
                                                                                                                                                                                                                           88.6%; Score 109; DB 2; Length 24; 95.7%; Pred. No. 2.1e-08; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                               HIV-1 vaccine synthetic peptide SEQ ID NO:99.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Col 71-72; 41pp; English
                                                                                                                                                                                                                                                                                   2 NTRKSERIORGPGRAFVTIGKIG 24
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                                                                                              Disclosure; Fig 3; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                               AAW98904 standard; peptide; 24
                    Sia CDY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-00462507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93US-00073378.
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(CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                 22; Conservative
                                         WPI; 1998-556461/47
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                     Klein MH,
                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                       Sequence 24 AA;
                                                                                                                                                                                                                                                                                                                                                                                          05-MAY-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                     AAW98904;
                                                                                                                                                                                                                           Query Match
                     Chong P,
                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence represents a fragment of a HIVI protein, and can be used in the immunogenic composition of the invention. The composition comprises a synthetic fusion polypeptide which includes a sequence encoding 1 or more I cell epitopes and a sequence encoding 1 or more B cell epitopes and a sequence encoding 1 or more B cell epitopes and a proteins. The composition are useful as vaccines against HIV infection. The composition induces HIV-1-specific polyclonal antibodies that are opsomiang and antiviral. The peptide components may be selected to induce a response against different viral isolates and in subjects who
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency virus, useful antigens in producing vaccines.
HIV infection and in diagnostic applications. AAW98892 to AAW98906, and AAW99899 to AAW9989 represent synthetic peptides from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV; vaccine; immunogenic composition; T cell epitope; B cell epitope; infection; antibody; antiviral.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
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Pred. No. 2.1e-08;
                                                                                                                                                                   88.6%; Score 109; DB 2; 95.7%; Pred. No. 2.1e-08;
                                                                                                                                                                                                                         0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY39769 standard; peptide; 24 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                         22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV1 chimeric peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chong P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-550482/46.
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                                                                                                                                                                                               Similarity
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                                                                                                                Sequence 24 AA;
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us-08-869-386-3.rag

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Enhancing immunogenicity of viral envelope glycoprotein - by co-
administration of viral envelope glycoprotein itself, and an oligopeptide
                                                                                                                                                                                                                                                                                                                                              A novel method of enhancing the immunogenicity of an envelope glycoprotein (EGP) of a virus (esp. HIV gpl20 or gpl60) in a host comprises admin. to the host at least one EGP of the virus in an amt. sufficient for priming vaccination and at least one peptide derived from an amino acid sequence of the EGP (e.g. the sequence shown), where the peptide comprises at least one virus-neutralisation epitope (VNE). The complex is able to enhance the induction of neutralising antibodies to the virus and to confer long lasting immunity, longer than 6 months. See also AAR36567-86. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV isolates HIV-IIIB and HIV-RF; hybrid peptide PR137; therapy; AIDS; principal neutralising domain; antibodies; diagnosis; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Grimaila R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus hybrid peptide RP137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Javaherian K, Farley J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 109; DB 2;
Pred. No. 2.1e-08;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                  Disclosure; Page 82; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 NTRKSERIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 NTRKSIRIQRGFGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR04475 standard; protein; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88US-00252949.
89US-00359543.
89US-00407663.
                                                                91US-00782154.
91US-00782241.
91US-00782252.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88US-00252949.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88.6%;
95.7%;
                                 92WO-EP002459
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Best Local Similarity 95.75
Watches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rusche JR, Putney SD,
Lynn DU, Petrobre J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
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                                                                                                                                       (INSP ) INST PASTEUR
                                                                                                                                                                                                         WPI; 1993-167398/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-SEP-2004
25-MAR-2003
20-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-OCT-1988;
                                                                                    28-OCT-1991;
28-OCT-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-OCT-1988;
                                 28-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-SEP-1989;
                                                                   28-OCT-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9003984-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-APR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUN-1989
13-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR04475;
                                                                                                                                                                        Girard M;
                                                                                                                                                                                                                                                                                   derive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This peptide is one example of an HIV-1 amplifier peptide for use in a composition for enhancing the immunogenicity of an envelope glycoprotein of a virus. The sequence corresponds to the major neutralisation epitope (loop v3) of HIV-1 brull isolate and enhances the induction of persistent neutralising antibodies in the host. The amplifier peptide is neutralising antibodies in the host. The amplifier peptide is neutralising antibodies. The compositions are particularly useful for vaccinating against HIV, SIV, HILV-I and HIV-II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Enhancing immunogenicity of envelope glyco:protein - for use as vaccine or immuno:therapeutic drug especially against HIV, HTLV-I and HTLV-II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus; gp120; gp160; EGP; VNE; immunity.
                                                                                                                                                                                                           human immunodeficiency virus; vaccine; human retrovirus; AIDS; acquired immunodeficiency syndrome; envelope glycoprotein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Virus neutralising epitope of envelope glycoprotein of HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 109; DB 2; Length 25;
Pred. No. 2.1e-08;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 NTRKSERIQRGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR36587 standard; peptide; 25 AA.
                                                                     AAR15058 standard; protein; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 13; Page 50; 71pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88.6%;
                                                                                                                                                                                                                                                                                                                                                                       90US-00494749
                                                                                                                                                                                                                                                                                                                                                                                                          90US-00494749.
                                                                                                                                                                          HIV-1 amplifier peptide #21
                                                                                                                                         (first entry)
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nes 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                           (INSP ) INST PASTEUR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1991-310366/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 25 AA;
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                                                                                                                                                                                                                                                                                                                                                                     19-MAR-1990;
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06-SEP-1993
                                                                                                                                         03-JAN-1992
                                                                                                                                                                                                                                                                                                   WO9114449-A
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                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Girard M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR36587;
                                                                                                        AAR15058;
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Matches

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Gaps

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Peptide RP137 comprises segments of the Principal Neutralising Domain (envelope protein) from isolates HIV-RP and HIV-IIIB. The last Cys residue is added for the purpose of crosslinking to carrier proteins. Cysteine residues may be added, so that the residues at or near both ends form a disulfide bond, giving peptide a loop-like configuration, which can be utilised to enhance immunogenic properties of the peptides. Protein is capable of eliciting, and/or binding with, neutralising antibodies. The neutralising domain is bounded by cysteine residues which occur at positions 296 and 331. The peptides can be used as immunogens or screening research or generate or identify poly. or monoclonal antibodies. See also AAR04427-R04505 and AAQ4273-Q04279. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monoclonal antibodies directed against regions of gpl20 of human immune deficiency virus-1 - are neutralising and able to inhibit infection of \mathtt{T}
                                                                                                                                                                                                                                                                                                                                                                    Revised record issued on 09-SEP-2004 : Correction to Feature Table Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Epitope, gpl20 protein, monoclonal antibody; HIV-1; antibody BAT123; antibody BAT267; antibody BAT1085, Toell infection inhibition; agultated immune deficiency syndrome; AIDS; AIDS-related complex; passive immunisation; antiviral; cytotoxic;
                               Principal neutralising domain of HIV variants - used for producing peptide(s) and antibodies for diagnosis, prophylaxis and/or therapy therapy of HIV infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    86.2%; Score 106; DB 2; Length 25;
87.5%; Pred. No. 5.7e-08;
tive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chang TW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chang NT,
                                                                                                  Claim 8 (58); Page 76; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 NNTRKSERIQRGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 NNTRKSIRITKGPGRAFVTIGKIG 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           viral load measurement; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Epitope of HIV-1 gp120 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW87618 standard; peptide; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87US-00057445.
87US-00137861.
91US-00767533.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 87.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sun BNC,
WPI; 1990-147824/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-095002/08
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                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-DEC-1987;
26-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAY-1987;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW87618;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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Gapa

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                                                                              The present sequence represents an epitope of the gp120 protein of human immune deficiency virus (HIV)-1. The sequence comprises amino acids 298 to 322 of gp120. The specification describes monoclonal antibodies which bind to sequences derived from the present epitope. Specifically, these antibodies are designated BATA13, 267 and 085. Monoclonal antibodies neutralise HIV-1, inhibiting both infection of T cells and formation of syncytia, so are used to treat acquired immune deficiency syndrome (AIDS) and AIDS-related complex, by passive immunisation, as carriers of cytotoxic or antiviral agents, and in extracorporeal systems. They can also be used as immunoassay reagents (for diagnosis or measurement of viral load) and to screen for neutralising epitopes, potentially useful in vaccine development. (Updated on 20-MAR-2003 to correct PR field.) (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antibodies able to neutralise more than one HIV-1 isolate are claimed. The gpl20 V3 loop sequences from different isolates comprising the Principal Neutralising Determinant motif GPGR are given in AAR42153-R42161. (Updated on 25-MRR-2003 to correct PN field.) (Updated on 24-OCT-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New recombinant human antibody - with HIV neutralising activity against
at least two isolates, useful for preventing or treating infection in
diagnosis, etc.
                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human Immunodeficiency Virus; antigen; ELISA; recombinant antibody; HIV-neutralising monoclonal antibody; immunoglobulin; AIDS; acquired immune deficiency syndrome; chimeric antibody; surface glycoprotein gpl20; V3 loop; epitope mapping.
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 used for treatment, prevention
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                                                                                                                                                                                                                                                                                                                                                                   85.4%; Score 105; DB 2; Length 25; 95.5%; Pred. No. 8e-08; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfarr DS;
cells and formation of syncytia, used for treatme
diagnosis of acquired immune deficiency syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus 1; (IIIB isolate)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gp120 v3 loop sequence of HIV-1 IIIB isolate.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 16; Page 100; 154pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR42153 standard; peptide; 22
                                                    Claim 2; Col 8; 16pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93WO-US002629.
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                                                                                                                                                                                                                                                                                                                                                                                                        21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1993-336600/42.
                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                    Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-APR-1992;
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25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR42153;
                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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This sequence represents the V3 loop from the gpl20 protein of the human minumundeficiency virus-1 (HIV-1) strain IIIB. The invention relates to the generation of heterohybridomas producing human monoclonal antibodies (see AAX79204-X79207) to a neutralising epitope of HIV-1 prepared by transforming peripheral blood lymphocytes with Spstein-Barr virus. The antibodies can be used to treat someone infected with HIV-1 or suffering from AIDS. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-idiotypic, antibody, gp120, HIV; human immunodeficiency virus, paratope; complemetarity determining region; CDR; immunisation; vaccine;
                                                                     Light chain; variable region; human; HIV-1; gpl20; monoclonal antibody; epitope; V3 loop; heterohybridoma; human immunodeficiency virus-1; peripheral blood lymphocyte; Epstein-Barr virus; EBV; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heterohybridoma producing human monoclonal antibodies to human immunodeficiency virus-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Binding site of BAT123 and BAT267 HIV antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         83.7%; Score 103; DB 2;
95.5%; Pred. No. 1.4e-07;
iive 0; Mismatches 1;
                                      HIV-1 strain IIIB gp120 V3 loop sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Simian-Human immunodeficiency virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Col 24; 42pp; English.
                                                                                                                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 paratope; complemetarity determ
immunotoxin; T-cell; AIDS; ARC.
                                                                                                                                                                                                                                                                                  90US-00538451.
91US-00684090.
92US-00872675.
                                                                                                                                                                                                                                                   94US-00345321.
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                                                                                                                                                                                                                                                                                                                                                                                          Gorny MK, Zolla-Pazner S;
 17-AUG-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 95.57
Local Similarity 95.57
Local 21; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-370481/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22 AA;
                                                                                                                                                                                                                                                   21-NOV-1994;
                                                                                                                                                                                                                                                                                  15-JUN-1990;
                                                                                                                                                                                                                                                                                                      12-APR-1991;
23-APR-1992;
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01-OCT-1991
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                                                                                                                                                                             US5914109-A.
                                                                                                                                                                                                                22-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is the sequence of the V3 loop of the gp120 envelop protein from HIV the sequence was used with a construct comprising part of the HIV-1 gp120 V3 loop common consensus PND sequence (AAM07399) fused to part of the HIV-1 CD4 binding site (AAW07391) and with the V3 loop part of the HIV-1 strains Thai B (AAW07393) or HGP-30 (AAW07394) to generate a lysine branched peptide which is useful for the prevention and treatment of AIDS. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AIDS vaccine - comprises branched lysine peptide fragments derived
                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                      HIV-1; gpl20; V3 loop; common consensus PND domain; envelop; CD4; binding site; stem-loop; lysine branched peptide; AIDS.
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                                                                     Score 103; DB 2; Length 22; Pred. No. 1.4e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       83.7%; Score 103; DB 2;
95.5%; Pred. No. 1.4e-07;
iive 0; Mismatches 1;
                                                                                    Pred. No. 1.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                     HIV-1 strain IIIB gp120 V3 loop sequence.
                                                                                                                                                             1 TRKSIRIQRGPGRAFVTIGKIG 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 5-6; 8pp; Japanese.
                                                                                                                                       3 TRKSERIORGPGRAFVTIGKIG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 NTRKSERIORGPGRAFVTIGKI 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NTRKSIRIORGPGRAFVTIGKI 22
                                                                                                                                                                                                                                                              AAW07392 standard; peptide; 22 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus 1.
2003 to standardise OS field)
                                                                   83.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95JP-00038835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95JP-00038835
                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                Query Match
Best Local Similarity 95.5
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from HIV env protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (TERU ) TERUMO CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-461278/46
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (OKUD/) OKUDA K.
                                  Sequence 22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-SEP-1996.
                                                                                                                                                                                                                                                                                                                                  16-OCT-2003
24-FEB-1997
                                                                                                                                                                                                                                                                                                  AAW07392;
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AAW07392
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Matches

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Gaps

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Example 10; Col 12; 15pp; English
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                    WPI; 1999-008810/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-DEC-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                   The peptide corresponds to residues 294-318 of the gpl20 envelope protein of HIV-1 which is a principal neutralising determinant (FND). Abs recognise residues 294-318 (MAb BAT267) or 304-318 (MAb 123). These MAb are used to raise anti-idiotypic Abs (AAbs). The Abs are useful for passive immunisation and as components for immunotoxins which destroy rells infected with HIV. They inhibit T-cell infection and syncytium formation, are group specific and neutralise specific strains of HIV-1. They can be used to treat AIDS or ARC. The AAbs can be used for active immunisation or can be admin with another vaccine to increase
                                                                                                                                                                Monoclonal antibodies specific to the gp120 HIV envelope protein - for immunisation against HIV in treatment of AIDS or ARC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV-1; gpl20; epitope; monoclonal antibody; envelope; neutralise; inhibit; infection; T-cell; inhibit syncytium formation; AIDS.
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                                                                                           Chang NT;
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                                                                                           Sun BNC,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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/label= peptide_a
11. .25
/label= peptide_b
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                                                                                                                                                                                                                       Claim 5; Page 97; 124pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus 1.
                                                                                         Sun CRY,
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87US-00137861.
89US-00343540.
92US-00895197.
89US-00454161.
90US-00531789.
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                                                     (TANO-) TANOX BIOSYSTEMS IN
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                         Chang TW, Fung MSC,
                                                                                                                           WPI; 1991-222664/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
nes 20, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (TANO-) TANOX
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21-DEC-1989;
12-JUN-1990;
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24-DEC-1987;
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05-JUN-1992;
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13-JAN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                              os field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Matches
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AAW72819
ID AAW72
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AC AAW72
DT 13-DA
DT 13-JA
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The present invention describes an antibody conjugate comprising an antibody (Ab) which binds to an epitope within amino acid residue 308-322 of gpl20 and neutralises HIV-1, conjugated with a cytotoxic agent, an anti-viral agent or an agent which facilitates passage through the blood brain barrier. Also described is an antibody conjugate as above but where the brids to an epitope within amino acid residue 398-312 of gpl2 which neutralises HIV-1. The present sequence represents an HIV-1 gpl20 epitope corresponding to positions 294 to 318. The Ab are monoclonal Ab which bind to the gpl20 protein on the envelope of HIV-1. They inhibit the infection of T-cells and also inhibit syncytium formation. The antibodies are group specific and neutralise different strains and isolates of HIV-1. The antibodies have a variety of uses, including the treatment and prevention of AIDS and AIDS related complex. They are especially used to kill infected T-cells. (Updated on 17-OCT-2003 to
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glycoprotein 140; gp140; neutralising antibody; conformational epitope;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Producing neutralizing antibodies useful for preventing, treating and diagnosing an HIV infection in a mammal comprises administering recombinant uncleaved gp140 proteins to a human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
Antibody conjugate comprising monoclonal antibody - which binds to epitope within amino acid residue of gpl20 which neutralises HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  81.3%; Score 100; DB 2; Length 25; 90.9%; Pred. No. 4.1e-07; ive . 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (USSH ) US DEPT HEALTH & HUMAN SERVICES.
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                                                                                     conjugated with, e.g. cytotoxic agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV-1 IIIB V3 loop peptide sequence.
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                                                                                                                                                              Disclosure; Col 8; 22pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93US-00165314.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20; Conservative
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loop peptide sequence. The peptide sequence is used to test the reactivity of the antibodies of the invention. The invention relates to a method for the production of neutralising antibodies against conformational epitopes of HIV-1 envelope proteins in humans. The method comprises administering to a human, a recombinant uncleaved gpl40 protein retaining its oligomeric structure. The human produces neutralising antibodies against conformational epitopes of the HIV-1 gpl40 protein found on the oligomeric structure of the gpl40. The anti-HIV-1 gpl40 antibodies of the invention can be used for preventing and disponsing an HIV infection in a mammal. Gpl40 antibodies are useful for treating an HIV infection. A disgnostic method using the antibodies involves isolating a body fluid, preferably blood, and contacting it with a labelled monoclonal antibody for gpl40, and detecting any bound antibody. (Updated on 12-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency virus infection.
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  sequence represents a human immunodeficiency virus type-1 IIIB V3-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #12.
                                                                                                                                                                                                                                                                                                                                                                                80.5%; Score 99; DB 3; Length 22; 95.2%; Pred. No. 5.1e-07; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 10; Page 119; 154pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 NTRKSERIQRGPGRAFVTIGK 22
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                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 95.2
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scott D, Zambidis E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI, 1998-506315/43
                                                                                                                                                                                                                                                                                                                                           Sequence 22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9836087-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW76842;
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Gaps

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optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, as prevents or inhibite 'bystander' apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral T cell immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The synthetic epitope is used in a complex with either the hepatitis B core antigen (HBc) or a sol. cleavage prod. of HBc (HBe), to prepare a vaccine. The peptide corresponds to residues 308-327 of the HIV env glycoprotein 120, with an N-terminal initiation Met. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus; retrovirus; vaccine; antibodies; HbC; HBe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gapa
                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antigen granule comprising HBC or HBE antigen - and HIV neutralised epitope obtd. by expression of recombinant prod., for e.g. vaccine.
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                                                                                                                                                                                 Length 20
                                                                                                                                                                                                               1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Epitope comprising residues 308-327 of HIV env gp 120.
                                                                                                                                                                               76.4%; Score 94; DB 2; 395.0%; Pred. No. 2.4e-06; ive 0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim Disclosure; Fig 4; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antigen; hepatitis B virus; HBV; core.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KAGA ) KAGAKU OYOBI KESSEI RYOHO.
                                                                                                                                                                                                                                                                                                                                                             AAR04060 standard; peptide; 21 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 NNTRKSERIQRGPGRAFVTI 20
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                                                                                                                                                                                                                                              2 NTRKSERIQRGPGRAFVTIG 21
                                                                                                                                                                                                                                                                            1 NTRKSIRIQRGPGRAFVTIG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88JP-00220770.
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                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                                                                                 Query Match
Best Local Similarity 95.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19; Conservative
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Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; AAQ02417.
                                                                                                                                                 Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JP02069194-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-SEP-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-SEP-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
23-JUL-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAR-1990.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                            AAR04060;
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RESULT 38

us-08-869-386-3.rag

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HIV-1 Btrain HXB2 gp120 V3 loop peptide amino acids 302-322.
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                                                                                                                                                    30-MAY-1995;
                                                                                                                                                                          15-JUN-1992;
                                                                                                                                                                                      09-JUN-1993;
                                                                                                       US5866137-A.
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                                                                                                                              02-FEB-1999
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                                                                                                                                                                                                                                    Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY16052;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      셤
                                                                                                                                                                                                                                                                                                                                                                                      AAR93071-R93074 represent sequences used as antigenic marker epitopes in a non-infectious retrovirus-like particle of the invention. This sequence crepresents the antigenic peptide CLTB3. The retrovirus-like particle contains 1-4 repeats of this sequence (or AAR93061). The coding sequence for the retroviral particle of the invention comprises a modified retroviral genome deficient in long terminal repeats, but containing the gas, pol and env genes in their natural genomic arrangement, along with the antigenic composition capable of eliciting a retroviral specific immune response. The composition is for parenteral or mucosal administration, preferably oral, anal, vaginal or intranásal administration. The specific immune response, such as against AIDS and AIDS related compositions. The particles may also be used in the prophylactic (or specific immune response, such as against AIDS and AIDS related conditions. The particles may also be used in the prophylactic (or curative) treatment of AIDS and related conditions, by acting to displace the binding of the HIV virus to human or animal cells, or by disrupting the 3-dimensional organisation of the virus. The particle can also be used to identify antibodies specifically reacting with retrovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                  Antigenically marked non-infectious retrovirus-like particles - used to vaccinate against, and in the treatment of, AIDS and AIDS related conditions.
                                                                                          Antigen; non-infectious; retrovirus; antigenic marker; immune response; long terminal repeat; gag; pol; env; AIDS; HIV; antibody; therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     76.4%; Score 94; DB 2; Length 21; 90.5%; Pred. No. 2.5e-06; iive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                       Klein MH;
                                                                                                                                                                                                                                                                       Persson R,
                                                                                                                                                                                                                                                                                                                                                                     Example 4; Page 38; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW75478 standard; peptide; 21 AA
AAR93073 standard; peptide; 21 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 NTRKSERIQRGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95WO-CA000483
                                                                                                                                                                                                                          94US-00290105
                                                                                                                                                                                                                                                                       Yao F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                                              27-SEP-1996 (first entry)
                                                                                                                                                                                                                                                (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19; Conservative
                                                                    Antigenic peptide CLTB73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
                                                                                                                                                                                                                                                                       Cao S,
                                                                                                                                                                                                                                                                                               WPI; 1996-139690/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 AA;
                                                                                                                                                                                                 15-AUG-1995;
                                                                                                                                                                                                                          15-AUG-1994;
                                                                                                                                                     WO9605292-A1
                                                                                                                                                                           22-FEB-1996.
                                                                                                                                                                                                                                                                       Rovinski B,
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27-APR-1999
                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW75478;
                       AAR93073
                                                                                          Antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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AAW75478
ID AAW75
XX
AC AAW75
DT 17-OC
DT 20-MA
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This sequence represents a peptide from the V3 loop of the gp120 protein from the human immunodeficiency virus type 1 (HIV-1) strain HXB2. The peptide is used to determine antibody responses after immunisation with a self-assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle. The retrovirus-like particle comprises a modified HIV genome devoid of long terminal repeats (LTRS) and contains a nucleotide sequence coding for a chimeric envelope glycoprotein. The chimeric envelope glycoprotein has the HIV-1 gp120 conserved region 2 and a second retroviral envelope amino acid sequence from a heterologous strain of HIV-1, HIV-2, HTLV-1 or HTLV-11 inserted into the first retroviral envelope amino acid sequence from a heterologous strain of HIV amino acid sequence (see AAWT5474-W7547). The novel retrovirus-like particle is useful in vaccines against HIV. (Updated on 20-MAR-2003 to correct PA field.) (Updated on 17-OCT-2003 to standardise OS field)
V3 loop; gp120 protein; HIV-1; retrovirus-like particle; genome; HIV-2; long terminal repeat; LTR; chimeric; envelope; glycoprotein; HTLV-1; HTLV-II; vaccine; human T-lymphotropic virus.
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chimeric envelope glycoprotein; HIV-1 gp120; conserved region 2; HIV-1;
HIV-2; HTLV-1; HTLV-II; vaccine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunogenic retrovirus-like particle - with chimeric HIV-1 envelope protein containing heterologous retroviral amino acid sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rovinski B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 4; Col 7-8; 12pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY16052 standard; peptide; 21 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 NTRKRIRIQRGPGRAFVTIGK 21
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                                                                                                                                                                   Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cao SX, Haynes J,
                                                                                                                                                                                                                                                                                                                                                                                                             95US-00453745.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92US-00839751
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Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
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Example 4; Col 9-10; 12pp; English.
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Human immunodeficiency virus 1.
                                                                            useful as vaccines against HIV
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                                                                                                                                                                                                                                                                                                                                                       92US-00839751
93US-00073526
                                                                                                                                                                                                                                                                                                                                             95US-00452503
                                                                                                                                                                                                                                                                                                                                                                      (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                           Conservative
                                                            WPI; 1999-357220/30
                                        (ROVI/) ROVINSKI B.
                                                                                                                                                                                     Local Similarity
                                                                                                                                                                      Sequence 21 AA;
               30-MAY-1995;
                              09-JUN-1993;
                                                                                                                                                                                                                                                                                                                                             30-MAY-1995;
                         .5-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                       15-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                            09-JUN-1993;
    15-JUN-1999.
                                                                                                                                                                                                                                                            20-MAR-2003
24-FEB-1999
                                                                                                                                                                                         19;
                                                                                                                                                                                                                                                                                                                        US5849475-A.
                                                                                                                                                                                                                                                                                                                                  15-DEC-1998
                                                                                                                                                                                                                                                  AAW85568;
                                                                                                                                                                                Query Match
                                                  Cao SX,
                                                                                                                                                                                           Matches
                                                                                                                                                                                                                              RESULT 41
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The present sequence represents a Human immunodeficiency virus type 1 derived peptide. The peptide is used in the immunosasay diagnostic kit of the invention. The specification describes an immunosasay diagnostic kit for detecting antibodies in a sample, which comprises an antigen consisting of a self-assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle encoded by a modified HIV genome that is devoid of long terminal repeats and contains a nucleotide sequence containing HIV-1 gpl20 conserved region 2 and a second amino acid sequence containing an envelope sequence of a heterologous strain of HIV-1, HIV-2, HILV-1 or HILV-11. (Updated on 20-MMR-2003 to correct PR field.)
                                                                            Immunoassay diagnostic kit for detecting antibodies - comprising chimeric retrovirus-like particles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Differentiating between infection by human immunodeficiency virus (HIV) and antiserum generated by immunization against HIV, comprises use of non-infectious, non-replicating HIV-like particle with heterologous, antigenic anchor sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Retrovirus-like particle CLTB73 containing a V3 (HXB2) antigenic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus; HIV; retroviral antigen; gag; pol; env; immune response; antigenic marker; antigenic epitope; retrovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 76.4%; Score 94; DB 2; Length 21; Best Local Similarity 90.5%; Pred. No. 2.5e-06; Matches 19; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Klein MH;
  Rovinski B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yao F, Persson R,
                                                                                                                                                    Example 4; Col 9-10; 12pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU08699 standard; peptide; 21 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 NTRKSERIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 NTRKRIRIORGPGRAFVTIGK 21
Cao SX, Haynes J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus.
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                                        WPI; 1999-069713/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rovinski B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JS6291157-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-SEP-2001.
  Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU08699;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 42
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                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes a nucleic acid molecule encoding a self assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle. The retroviral particle comprises a modified HIV genome devoid of long terminal repeats containing a nucleotide sequence coding for a chimeric envelope glycoprotein which has a first (a) and second (b) retroviral envelope amino acid sequence, where (a) contains the HIV-1 gpl20 conserved region 2, and (b) contains a retroviral envelope amino acid sequence of a hererologous strain of HIV-1, HIV-2, HIV-1 or HILV-11 inserted into (a) at an endogenous site (BglII and StuI). (b) may comprise peptides AAY16049-51 and AAY16055. The nucleic acids are useful as vaccines against HIV. The present sequence is used in the course of the invention. (Updated on 20-MAR-2003 to correct PR field.) (Updated on 17-0CT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunoassay diagnostic kit; antibody detection; chimeric envelope protein; HIV-1 gpl20 conserved region 2; HIV-1; HIV-2; HTLV-1; HTLV-1.
                                                                                                                                                                                                                                                                                                    Immunogenic retrovirus like particles comprising modified HIV genomes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  76.4%; Score 94; DB 2; Length 21; 90.5%; Pred. No. 2.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus type 1 derived peptide.
                                                                                                                                                                                                                 Klein MH, Haynes J, Rovinski B;
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                                                              95US-00452520
                                                                                                     92US-00839751
93US-00073526
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(first entry)
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The invention relates to a method for determining the presence of antibodies specifically reactive with HIV retroviral antigens in a sample. This involves contacting a sample suspected of containing HIV-

Disclosure; Col 17; 28pp; English.

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Gaps ö

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specific antibodies with a non-infectious, non-replicating, immunogenic HIV-like particle as an antigen. The antigen comprises an assembly of a gag gene product. A pol gene product and a modified env gene product containing a non-retroviral heterologous, antigenic, anchor sequence that replaces the endogenous anchoring functions of the env gene product. The method detects immune complex formation between HIV-specific antibodies and the antigens. The method is also useful for identifying antiserum generated by immunisation with an immunogenic composition capable of alliciting HIV-specific immune response. The antigenic marker may comprise at least one antigenic epitope from another virus. This sequence represents a retrovirus-like particle containing an antigenic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A genetic vaccine against HIV contains a DNA construct which comprises the sequence encoding gp160. The genetic material was then introduced into the cells of an individual by (a) contacting the individual's cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunisation; vaccine; therapy; prophylaxis; defective gene;
non-functional gene; template; antisense; ribozyme; bupivacaine;
human immunodeficiency virus; acquired immune deficiency syndrome; HIV;
                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Method for introducing genetic material into cells - utilises polynucleotide function enhancer and nucleic acid free of retroviral particles, e.g. HIV immunisation.
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                                                                                                                                                                                                                'Match 76.4%; Score 94; DB 4; Length 21; Local Similarity 90.5%; Pred. No. 2.5e-06; les 19; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Merva MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Coney LR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 44; 136pp; English.
                                                                                                                                                                                                                                                                             2 NTRKSERIQRGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                       AAR57470 standard; protein; 22 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wang B,
                                                                                                                                                                                                                                                                                             1 NTRKRIRIQRGPGRAFVTIGK 21
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93US-00029336.
93US-00093235.
93US-00124962.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV BRU V3 loop peptide.
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WILLIAMS W V.
WANG B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZURAWSKI V R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-263787/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CONEY L R.
MERVA M J.
                                                                                                                                                                                     Sequence 21 AA;
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21-SEP-1993;
21-SEP-1993;
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11-MAR-1993;
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21-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-AUG-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                     AAR57470;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MEIN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MERV/)
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                                                                                                                                                                                                                                               Matches
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with a polymucleotide function enhancer (bupivacaine) and (b) administering to the cells the nucleic acid molecule free of retroviral particles. Nucleic acid molecules which are delivered to cells may serve as genetic templates for proteins that function as prophylactic and/or therapeutic immunising agents; replacement copies of defective, missing or non-functional genes; genetic templates for therapeutic proteins; genetic templates for antisense molecules or as genetic templates for ribozymes. This peptide was derived from the V3 loop of an HIV strain (an epicope targetted by HIV neutralising antibodies) and was used to ederemine whether the anti-gp160 antibodies elicited in mice immunised with the genetic vaccine were reactive with this region. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a composition comprising activated alpha-2-macroglobulin (alpha_2M asterisk ), 3-0-deacylated monophosphoryl lipid A (mpL) and granulocyte macrophage colony stimulating factor (GM-CSF). The invention also relates to an adjuvant suitable for use in multivalent HIV immunogenic compositions. The compositions is useful for eliciting an type 1 (HIV-1) v3IIIB peptide used in the exemplification of the invention. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1; HIV-1; adjuvant; immunomodulator; alpha-2-macroglobulin; 3-0-deacylated monophosphoryl lipidA; MPL; GM-CSF; granulocyte macrophage colony stimulating factor; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of 2-macroglobulin (2Masterisk), 3-O-deacylated monophosphoryl lipid A (MPL) and granulocyte macrophage colony stimulating factor (GM-CSF) for
                                                                                                                                                                                                                                                                                                                                      Gapa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus type 1 (HIV-1) V3IIIB peptide.
                                                                                                                                                                                                                                                                                              76.4%; Score 94; DB 2; Length 22; 90.5%; Pred. No. 2.6e-06; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 21; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                            2 NTRKSERIQRGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                  2 NTRKRIRIQRGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAE20149 standard; peptide; 24 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-AUG-2001; 2001WO-US026589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-AUG-2000; 2000US-0227624P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eliciting an immune response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 90.5
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vaccine, V3IIIB peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Haynes BF, Liao H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-269315/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYDU-) UNIV DUKE.
                                                                                                                                                                                                                                                           Sequence 22 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New peptide(s) comprising HIV gpl20 epitope(s) - for prodn. of vaccines against HIV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus type 1; HIV-1; gpl20 epitopes; vaccines; HIV neutralising antibodies.
                                                                              Gaps
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Query Match 76.4%; Score 94; DB 5; Length 24; Best Local Similarity 95.0%; Pred. No. 2.9e-06; Matches 19; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jeansson S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV-1 gp120-23 epitope amino acids 296-230.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Svennerholm B, Rymo L,
                                                                                                                                                                                                                                                                                                                                                                                                AAR63820 standard; peptide; 25 AA.
                                                                                                                                             1 NNTRKSERIORGPGRAFVTI 20
                                                                                                                                                                                      5 NNTRKSIRIQRGPGRAFVTI 24
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25-MAR-2003 (revised)
29-JUN-1995 (first entry)
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Search completed: May 16, 2005, 13:05:13 Job time : 171.846 secs

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GenCore version 5.1.6
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- protein search, using sw model OM protein Run on:

May 16, 2005, 12:39:11 ; Search time 137.231 Seconds (without alignments) 89.556 Million cell updates/sec

US-08-869-386-3 123 1 NNTRKSERIQRGPGRAFVTIGKIG 24 Title: Perfect score:

Sequence:

**BLOSUM62** Scoring table: 1612378 segs, 512079187 residues Searched:

Gapop 10.0 , Gapext 0.5

16988 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

UniProt\_03:\*
1: uniprot\_sprot:\*
2: uniprot\_trembl:\* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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, ,		<b>09e887</b>	08aqx9	Q8aqy0	010481	09gex7	QBagy	28ady2	07zjt3	09pxf1	Q9enm9	078324	078326	078345	078378	078381	Qetgte	078328	Q78323	<b>Q78327</b>	<b>Q78380</b>	Q6u2m7	09zg65	gguck9	Q9uc10	Q16228	Q6eml0	Q6eml1	Q924c7	094781	011890	011891
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§ F	3	Q9E8S7	OBAQX9	OBAQYO	010481	090EX	Q8AQY:	QBAQY2	Q7ZJT3	O9PXF1	O9ENM9	078324	078326	078345	078378	078381	QETQT6	078328	Q78323	078327	078380	Q6U2M7	09ZG65	Q9UCK9	OBUCLO	016228	Q6EML0	Q6EML1	Q924C7	094781	011890	011891
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Pred. No. 0.0061;
2; Mismatches 5; Indels
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NON TER 23 23
SEQUENCE 23 AA; 2596 MW; 6C038F27BC0CA1E0 CRC64;
                                                                                                                                 01-WAR-2001 (TrEMBLrel. 16, Created)
01-WAR-2001 (TrEMBLrel. 16, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Envelope glycoprotein (Fragment).
                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
                                                                  23 A.A.
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                                                                  PRT;
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1 Similarity 60.9%;
14; Conservative 2
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Q8AQX9
Q8AQX9;
Q8AQX9;
Q1-MAR-2003 (TrEWBLrel. 23, Created)
01-MAR-2003 (TrEWBLrel. 23, Last sequence update)
01-OCT-2003 (TrEWBLrel. 25, Last annotation update)
Envelope glycoprotein (Fragment) Human immunodeficiency virus 1. Viruses; Retroid viruses; Retroviridae; Lentivirus. NCBI\_TaxID=11676; REAR REAR SOCIAL DESTRUCTION OF THE PROPERTY O

SEQUENCE FROM N.A.
BEDLINE=22860939; PubMed=14502005;
Freel S., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,
Patrick E., Lennox J.L., Hicks C.B., Eron J.J. Jr., Shugars D.C.,

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MEDLINE-21103026; PubMed=11170057;

MEDLINE-21103026; PubMed=11170057;

MEDLINE-21103026; PubMed=11170057;

MEDLINE-21103026; PubMed=11170057;

MEDLINE-21103026; PubMed=11170057;

Lin H.J., Siwak E.B., Lauder I.J., Hollinger F.B.;

Lin H.J., Siwak E.B., Lauder I.J., Hollinger F.B.;

"Long-term culture of human immunodeficiency virus type 1 resulting in 10ss of glycosylation sites.";

J. Med. Virol. 63:197-202(2001).

M. Med. Virol. 63:197-202(2001).

M. Med. Virol. 63:197-202(2001).

M. GO; GO:0015021; C:integral to membrane; IEA.

GO; GO:0015021; C:integral to membrane; IEA.

GO; GO:0015098; F:structural molecule activity; IEA.

GO; GO:0019031; C:viral envelope; IEA.

GO; GO:0019031; C:viral envelope; IEA.

M. GO; GO:001505; F:structural molecule activity; IEA.

M. MINEPRO; IPR001056; Pept_S24_S26_C.

M. Pfam; PF00516; GP120; 1.

M. AIDS; Coat protein; Envelope protein; Glycoprotein; Transmembrane.
                                                                           SEQUENCE FROM N.A.
MEDLINE=97255649; PubMed=9100996;
MENCHEN S.D., Lockey T.D., Slobod K.S., Hurwitz J.L.;
"Drift from the GPGRAF HIV-1 envelope V3 crown sequence in a North American inner city.",
AIDS Res. Hum. Retroviruses 13:527-528(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Coat protein; Envelope protein; Glycoprotein; Transmembrane.
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25 AA; 2801 MW; 25E1B150CD7C14B6 CRC64;
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O1.MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1.JUN-2003 (TrEMBLrel. 24, Last annotation update)
Viruses, Retroid viruses, Retroviridae, Lentivirus.
NCBL_TaxID=11676;
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
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Pred. No. 1.5;
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61.9%; Pred. No. 0.33;
:ive 0; Mismatches
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tes 13, Conservative
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Rreel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,
Rreel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,
Rreel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,
Rreel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,
R. Patrick B., Lennox J.L., Hicks C.B., Bron J.J. J. Shugars D.C.;
R. "Envelope diversity, coreceptor usage and syncytium-inducing phenotype
R.T. of HIV-1 variants in saliva and blood during primary infection.";
R. MIDS 17:2025-2033(2003).
R. GO; GO:0016021; C:integral to membrane; IEA.
R. GO; GO:0019028; C:viral capaid; IEA.
R. GO; GO:0019028; C:viral envelope; IEA.
R. GO; GO:0019028; C:viral envelope; IEA.
R. GO; GO:0019031; C:viral envelope; IEA.
R. R. InterPro: IPR00777; GP120.
R. BIVELOPE protein.
R. NON-TER.
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         "Envelope diversity, coreceptor usage and syncytium-inducing phenotype of HIV-1 variants in saliva and blood during primary infection."; AIDS 17:2025-2033(2003).
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NON TER 25 25

SEQÜENCE 25 AA; 2749 MW; 9B6E9DACB8D56C0C CRC64;
                                                                                             EMBL, AF536914; AAN63929.1; -. GO, GO:0016021; C:integral to membrane; IEA. GO; GO:0019028; C:viral capsid; IEA. GO; GO:0019031; C:viral capsid; IEA. GO; GO:0019031; C:viral envelope; IEA. GO; GO:0005198; F:structural molecule activity; IEA. Integro; IPR000777; GP120.
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
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58.3%; Pred. No. 0.069;
trive 1; Mismatches
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|NNTRRS--INIGPGRAFYATDIIG 23
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1es 14; Conservative
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7 NNTRRS--IPLGOGRAWFITG 25

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Human immunodeficiency virus 1.

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Preel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,

Patrick E., Lennox J.L., Hicks C.B., Eron J.J. Jr., Shugars D.C.;

Envelope diversity, coreceptor usage and syncytium-inducing phenotype

of HIV-1 variants in saliva and blood during primary infection.";

AIDS 17:2025-2033(2003).

EMBL, PR536912; ANG327.1; --

GO, GO:0019031; C:viral envelope; IEA.
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Freel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,
Patrick E., Lennox J.L., Hicks C.B., Eron J.J. Jr., Shugars D.C.;
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of HIV-1 variants in saliva and blood during primary infection.";
AIDS 17:2025-2033 (2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 / Match 37.4%; Score 46; DB 2; Length 25; Local Similarity 54.2%; Pred. No. 3.4; Pred. 13; Conservative 0; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9; Indels
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25 AA; 2601 MW; 71B5A774CE256C09 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25 AA; 2601 MW; 71B5A774CE256C09 CRC64;
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
                                                                                                                     01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
                                                                                                                                                                                                                                                                                      Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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                                                                                                                                                                                                                   Envelope glycoprotein (Fragment).
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                                                            PRELIMINARY;
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MEDLINE=22439926; PubMed=12552446;

A lversen A.K.N., Christlansen C.B., Attermann J., Eugen-Olsen J.,

Schulman S., Berntorp E., Ingeralev J., Fugger L., Schaibel E.,

Schulman S., Berntorp E., Ingeralev J., Fugger L., Schaibel E.,

Tengborn L., Gerstoff J., Dickmeiss E., Svejgaard A., Skinhoj P.;

Tengborn L., Gerstoff J., Dickmeiss E., Svejgaard A., Skinhoj P.;

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Tengborn L., Gerstoff J., Dickmeiss E., Svejgaard A., Skinhoj P.;

Tuntert J. Infect. Dis 187:125-225(2003).

In J. Infect. Dis 187:125-225(2003).

R GO; GO:0015021; C:integral to membrane; IEA.

GO; GO:0015021; C:integral to membrane; IEA.

GO; GO:0015021; C:viral capsid; IEA.

GO; GO:0015021; C:viral envelope; IEA.

RO; GO:0015021; C:viral envelope; IEA.

FinterPro; IPR000777; GP120.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Coat protein; Envelope protein; Glycoprotein; Transmembrane
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
                                                                                                                                  Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBL_TaxID=11676;
                            (TrEMBLrel. 24, Last sequence update) (TrEMBLrel. 26, Last annotation update)
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01-JUN-2003 (TrEMBLrel. 24, Created)
                                                                                           Envelope glycoprotein (Fragment).
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Guery Match

Best Local Similarity 64.37
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RESULT 10

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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Immunodeficiency virus type 1, viral sample FLPARSC (Florida patient A), partial env cds. V5 region. (Fragment).
Human immunodeficiency virus 1.
Viruses, Retroid viruses; Retroviridae, Lentivirus.
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01-OCT-2002 (TrEMBLrell. 22, Last annotation update)
Immunodeficiency virus type 1, viral sample FLPARSF (Florida patient A), partial env cds, V5 region. (Fragment).
Human immunodeficiency virus 1.
Viruses, Retroid viruses; Retroviridae, Lentivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Molecular epidemiology of HIV transmission in a dental practice."; Science 256:1165-1171(1992).
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Science 256:1165-1171(1992)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=92271245; PubMed=1589796;
O. C.-Y., Ciestelski C.A., Myers G., Bandea C.I., Luo C.C.,
Korber B.T.M., Mullins J.I., Schocherman G., Berkelman R.L.,
Economou A.N., Witte J.J., Furman L.J., Satten G.A., Curran J.W.,
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Ou C.-Y., Ciesielski C.A., Myers G., Bandea C.I., Luo C.C.,
Korber B.T.M., Mullins J.I., Schochetman G., Berkelman R.L.,
Economou A.N., Witte J.J., Furman L.J., Satten G.A., Curran J.W.,
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Pred. No. 3.5e+02;
1; Mismatches '6; Indels
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Zhang L.Q., Leigh-Brown A.J.;
Zhang L.Q., Leigh-Brown A.J.;
Submitted (RFR-1992) to the EMBL/GenBank/DDBJ databases.
EMBL; M92111; AAA44467.1; -
NON TER 1 1 1
NON TER 1 1 1
SEQUENCE 17 AA; 1625 MW; 3283A0BFD3FCA370 CRC64;
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Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                17 AA
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NON TER 1

NON_TER 17 17

SEQUENCE 17 AA; 1635 MW;
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3 NNTKGSETFRPGGG 16
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                                                                                                                                                                                                                PRELIMINARY;
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01-NOV-1996 (TrEMBLrel.
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Q78326
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Q78345
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01-OCT-2002 (TEMBLrel. 22, Last annotation update)
Immunodeficiency virus type 1, viral sample FLPARSB (Florida patient A), partial env cds, V5 region. (Fragment).
Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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MEDLINE=92271245; PubMed=1589796;
Ou C.-Y., Ciesielski C.A., Myers G., Bandea C.I., Luo C.C.,
Korber B.T.M., Mullins J.I., Schochetman G., Berkelman R.L.,
ECONOMOU A.N., Witte J.J., Furman L.J., Satten G.A., Curran J.W.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
MEDLINE=20411423; PubMed=10954550;
DOI=10.11284/VI.74.18.8494-8501.2000;
Nelson J.A.E., Baribaud F., Edwards T., Swanstrom R.;
"Patterns of changes in human immunodeficiency virus type 1 "sequence populations late in infection.";
J. Virol. 74:8494-8501(2000).
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Pred. No. 2.46+02;
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Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases
EMBL; M92110; AAA44466.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23 23 23 23 23 2460 MW; 6108EAEC9C0CA947 CRC64;
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                                                                                                                                                                                                         (TrEMBLrel. 16, Created)
(TrEMBLrel. 16, Last sequence update)
(TrEMBLrel. 26, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                  Viruses, Retroid viruses, Retroviridae, Lentivirus, NCBI_TaxID=11676;
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                                                                                                                                          23 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            J. Virol. 74:8494-8501(2000).
EMBL; AF092639; AAD04382.1; -.
GO; GO:0019031; C:viral envelope; IEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 NNTRKSERIORGPGRAFVTIGKI 23
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                                                                                                                                                                                                                                                                                                                      Envelope glycoprotein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31.7%;
47.8%;
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Best Local Similarity 50.00,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 47.8 es 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
                                                                                                                                          PRELIMINARY;
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                                                                                                                                                                                                                                                     01-MAR-2001
01-MAR-2004
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SEQUENCE
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Q78324;
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RESULT 11

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Gaps

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1 NNTRKSERIQRGPG 14

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Gaps

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Score 31, DB 2, Length 17;
Pred. No. 7.6e+02;
2; Mismatches 6; Indels
                                      347570D2D12CA370 CRC64;
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                                                                                                                                                                                                24 AA
                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae (Baker's yeast).
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                                                                                                                                                                                                 PRT;
EMBL, M92126; AAA44496.1; -.
NON TER 1 1 1
NON TER 17 17
SEQÜENCE 17 AA; 1708 MW:
                                                                                                                                                                                                                                                                                                                                                                                                                                           25.2%;
                                                              25.2%;
                                                                                                              1 NNTRKSERIQRGPG 14
                                                                                                                                      3 NNTNNTETFRPGGG 16
                                                                                      6; Conservative
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                                                                                                                                                                                                PRELIMINARY;
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Best Local Similarity
Matches 6; Conserv
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05-JUL-2004
                                                                                                                                                                                                                                                                      Name=YHR065C
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Q6TQT6
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Immunodeficiency virus type 1, viral sample FLPBR5A (Florida patient B), partial env cds, VS region. (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 26, Last annotation update)
Immunodeficiency virus type 1, viral sample FLPBR5F (Florida patient B), partial env cds, V5 region. (Fragment).
                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
MEDIJINE=92271245; PubMed=1589796;
Ou C.-Y., Ciesielski C.A., Myers G., Bandea C.I., Luo C.C.,
Corber B.T.M., Mullins J.I., Schochetman G., Berkelman R.L.,
Economou A.N., Witte J.J., Furman L.J., Satten G.A., Curran J.W.,
Jaffe H.W.,
"Molecular epidemiology of HIV transmission in a dental practice.";
Science 256:1165-1171(1992).
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MEDLINE=92271245; PubMed=1589796;
MEDLINE=92271245; PubMed=1589796;
Ou C.-Y., Cieslelski C.A., Myers G., Bandea C.I., Luo C.C.,
Korber B.T.M., Mullins J.I., Schochetman G., Berkelman R.L.,
Economou A.N., Witte J.J., Furman L.J., Satten G.A., Curran J.W.,
Jaffe H.W.;
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      Length 17;
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42.9%; Pred. No. 5.2e+02;
tive 3; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                           Zhang L.Q., Leigh-Brown A.J.;
Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases
EMBL; M92123; AAA44493.1;
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Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 AA; 1723 MW; 34757935D12CA370 CRC64;
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NCBI_TaxID=11676,
                                                                                                                                                                                                                               Viruses, Retroid viruses, Retroviridae, Lentivirus, NCBI_TaxID=11676;
   26.0%; Score 32; DB 2; I
50.0%; Pred. No. 5.2e+02;
iive 1; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 AA.
                                                                                                                                       PRT;
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                                                      1 NNTRKSERIQRGPG 14
                                                                            3 NNTNGSETFRPGGG 16
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Matches 6; Conservative
                               Conservative
                                                                                                                                       PRELIMINARY;
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             Best Local Similarity
Matches 7; Conser
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      Query Match
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Q78378;
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Q78381
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Immunodeficiency virus type 1, viral sample FLPARSE (Florida patient A), partial env cds, V5 region. (Fragment).
Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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MEDLINE=92271245; PubMed=1589796;
Ou C.-Y., Ciestelski C.A., Myers G., Bandea C.I., Luo C.C.,
Korber B.T.M., Mullins J.I., Schochetman G., Berkelman R.L.,
Bconomou A.N., Witte J.J., Furman L.J., Satten G.A., Curran J.W.,
Jaffe H.W.;
                                                                                                                                                                                            Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetales; Saccharomycetales; Saccharomyces.
NCBI TaxID=4932;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
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Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases.
EMBL; M92113; AA44465.1; -.
NON TER 17 17
                                                                                                                                                                                                                                                                                                                                          STRAIN=AB972;
Kennedy M.C., Dietrich F.S.;
Submitted (EEP-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AY389302; AAQ97234.1; --
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24 AA; 2866 MW; 83820AB41EF59E7C CRC64;
05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
NAMESCOP (Fragment).
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Pred. No. 1.1e+03;
1; Mismatches 2;
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17 AA;

SEQUENCE

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PRELIMINARY;

078323

RESULT 18

SEQUENCE FROM N.A.

A PAC A PAC

NCBI\_TaxID=11676;

[2] SEQUENCE FROM N.A.

Jaffe H.W.;

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Gaps
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NAR-2004 (TrEMBLrel. 26, Last annotation update)
Immunodeficiency virus type 1, viral sample FLPBRSE (Florida patient B), partial env cds, VS region. (Fragment).
Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=92271245; PubMed=1589796;
U.C.-Y., Ciestellski C.A., Myers G., Bandea C.I., Luo C.C.,
Korber B.T.M., Mullins J.I., Schochetman G., Berkelman R.L.,
Economou A.N., Witte J.J., Furman L.J., Satten G.A., Curran J.W.,
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Ayre B.G., Blair J.E., Turgeon R.;
                                                                                                                                      Score 29; DB 2; Length 17;
Pred. No. 1.7e+03;
2; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 29; DB 2; Length 17;
Pred. No. 1.7e+03;
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Zhang L.Q., Leigh-Brown A.J.;
Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases
EMBL; M92112; AAA44468.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zhang L.Q., Leigh-Brown A.J.;
Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases.
EMBL; M92125; AAA44495.1; -.
                                                                           17 17
17 AA; 1651 MW; 34757BBFD12CA370 CRC64;
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CGUZM7;
05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Galactinol synthase (EC 2.4.1.123) (Fragment).
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                                                                                                                                        Query Match 23.6%;
Best Local Similarity 42.9%;
Matches 6; Conservative
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nes 6; Conservative
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SEQUENCE
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Q78380
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                                                                             Gaps
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Immunodeficiency virus type 1, viral sample FLPAR5A (Florida patient A), partial env cds, VS region. (Fragment).
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 02, Last sequence update)
Immunodeficiency wirus type 1, viral sample FLPARSD (Florida patient A), partial env cds, VS region. (Fragment).
Wiruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jaffe H.W.; "Molecular epidemiology of HIV transmission in a dental practice."; Science 256:1165-1171(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Molecular epidemiology of HIV transmission in a dental practice."; Science 256:1165-1171(1992).
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MEDLINE=92271245; PubMed=1589796;
Ou C.-Y., Ciesielski C.A., Myers G., Bandea C.I., Luo C.C.,
Korber B.T.M., Mullins J.I., Schochetman G., Berkelman R.L.,
Economou A.N., Witte J.J., Furman L.J., Satten G.A., Curran J.W.,
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                                       Length 17;
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Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases
EMBL; M92109; AAA44465.1; -.
NON TER 1
1750 MW; 346FD0BB802CA370 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17 AA; 1649 MW; 3E857BBFD12CA370 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                         Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                   Query Match 24.8%; Score 30.5; DB 2; Best Local Similarity 53.3%; Pred. No. 9.3e+02; Matches 8; Conservative 2; Mismatches 4;
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Pred. No. 1.7e+03;
2; Mismatches 6;
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23.6%;

Local Similarity 42.9

Best Loc Matches

SEQUENCE

Query Match

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PRELIMINARY;

078327

NCBI\_TaxID=11676;

SEQUENCE FROM N.A

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InterPro; IPR000096; Serum_amyloid_A. Pfam; PF00277; SAA; 1. SEQUENCE 16 AA; 1612 MW; 1CAB4F07
                                                                                                                     Query Match 22.0%;
Best Local Similarity 71.4%;
Matches 5; Conservative
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Matches 5, Conservative
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01-NOV-1996 (TrEMBLrel.
01-NOV-1998 (TrEMBLrel.
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1 RGPGGAW 7
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
"Functional and phylogenetic analyses of a conserved regulatory program in the phloem of minor veins."; Plant Physiol. 133:1229-1239(2003). EMBL; AX379782; A&Q74884.1; -. GO; GO:0047216; F:inositol 3-alpha-galactosyltransferase acti. . GO; GO:0016757; F:transferase activity, transferring glycosyl. . Glycosyltransferase; Transferase.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-WAR-2004 (TrEMBLrel. 26, Last annotation update)
Serum amyloid A isotype 2 alpha protein (Serum amyloid A protein)
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Chlamydia trachomatis.
Bacteria; Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia.
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Pred. No. 2.6e+03;
2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                    Length 22;
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STRAIN=L2 4348;
Wang L., Steenburg S.D., Zheng Y., Larsen S.H.;
Submitted (400-1998) to the EMBL/GenBank/DDBJ databases.
EMBL, AF087291; AAD04068.1; -
                                                                                                                                                                                                                         NON TER 22 22
SEQÜENCE 22 AA; 2295 MW; A6673B5BFD06430C CRC64;
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18 AA; 2026 MW; CB911767583AF4E3 CRC64;
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Pred. No. 2.2e+03;
0; Mismatches 1;
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85.7%;
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16 PKRAFVT 22
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SEQUENCE
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                                                                                               Gaps
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 3.4e+03;
1; Mismatches 1; Indels
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Pred. No. 3.4e+03;
1; Mismatches 1; Indels
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1CAB4F077C9C8CC1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         an individual patient of AA-amyloidosis.";
Biochim. Biochys. Acta 1180:195-200(1992).
GO; GO:0005576; C:extracellular; IEA.
GO; GO:0006953; P:acute-phase response; IEA.
Interpro; IPR000096; Serum_amyloid_A.
Pfem; PF00277; SAA; 1.
                                                                                                                                                                                                                                                                                                                                 16 AA.
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71.4%;
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Trypanosoma cruzi.
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae; Trypanosoma.
NCBI_TaxID=5693;
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"Glucagon-like peptide (GLP)-2 action in the murine central nervous system is enhanced by elimination of GLP-1 receptor signaling.";
Biol., Chem. 276:21489-21499 (2001).
EMBL; AF338224; AAK63043.1;
MGD; MGI-2135733; GLPZ.
GO; GO:0016021; C:integral to membrane; TAS.
GO; GO:0004967; F:glucagon receptor activity; TAS.
GO; GO:0007186; P:Glucagon receptor activity; TAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota, Meracasa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
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                                                                                                                                                                                 Length 19;
                                                                                                                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 27; DB 2; Length 22; Pred. No. 4.8e+03; 3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tanaka T., Tanaka M.;
Submitted (AUG-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; D87227; BAA133181; -.
NON TER.
                1 1
19 19
19 AA; 1985 MW; 79B9BEI06D048680 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               valuer-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUM-2003 (TrEMBLrel. 24, Last annotation update)
Glucagon-like peptide-2 receptor (Fragment).
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Last annotation update)
                                                                                                                                                                         Score 27; DB 2; I
Pred. No. 4.1e+03;
2; Mismatches 3;
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22 AA; 2526 MW;
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                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                        Query Match
Best Local Similarity
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01-DEC-2001 (
01-DEC-2001 (
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SEQUENCE
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Pubmed=15140948; DOI=10.1093/molbev/mshl57;

Axelsson E., Smith N.G., Sundstrom H., Berlin S., Ellegren H.;

Axelsson E., Smith N.G., Sundstrom H., Berlin S., Ellegren H.;

"Male-biased mutation rate and divergence in autosomal, z-linked and

w-linked introns of chicken and Turkey.";

Mol. Biol. Evol. 21:1538-1547(2004).

EMBL, AX139863; AAAV4867.1;

EMBL, AX139863; AAAV4867.1;

EMBL, AX139863; Fitransferase activity, IEA.

GO; GO:0016772; Fitransferase activity, transferring phosphor. . .; IEA.

InterPro; IRR000749; ATP-gua_Ptrans.

Pfam; PF02807; ATP-gua_Ptrans.
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PubMed=15140948; DOI=10.1093/molbev/msh157;
PubMed=15140948; DOI=10.1093/molbev/msh157;
Axelseon E., Smith N.G., Sundertrom H., Berlin S., Ellegren H.,
"Male-biased mutation rate and divergence in autosomal, z-linked and
w-linked introns of chicken and Turkey.";
wol. Biol. Evol. 21:1538-1547(2004).
EMBL; AX198962; AAN74866.1; --
GO, GO:0016301; F:kinase activity; IEA.
GO, GO:0016301; F:transferase activity, transferring phosphor. . .; IEA.
InterPro; IPR000749; ATP-gua_Ptrans.
Ffam; PF02807; ATP-gua_PtransN; 1.
                                                                                              Gaps
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Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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Meleagris gallopavo (Common turkey).

Meleagris gallopavo (Common turkey).

Meleagris metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Meleagris.
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        Score 27; DB 2; Length 17; Pred. No. 3.6e+03;
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Pred. No. 4.1e+03;
2; Mismatches 3; Indels
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25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
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47.1%; Pred. No. 4...
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22.0%;
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                                                                                      6; Conservative
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                                         Best Local Similarity
Matches 6; Conserv
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        Query Match
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06EML0
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AC 06EML
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"High prevalence of GB virus C strains genetically related to strains with Asian origin in Nicaraguan hemophiliacs.";
With Asian origin in Nicaraguan hemophiliacs.";
J. Med. Virol. 15149-155 (1997).
EMBL; U86117; AABSS851.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
MEDLINE=9732371; PubMed=9179760;
DOI=10.1002/(SICI)1096-9071(199706)52:2<149::AID-JMV5>3.3.CO;2-N;
Gonzalez-Perez M.A., Norder H., Bergstrom A., Lopez E., V1sona K.A.,
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                                                                                                    011893 PRELIMINARY; PRT; 25 AA.
011893;
01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-JUL-1997 (TrEMBLrel. 19, Last annotation update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Polyprotein (Fragment)
GB virus (Appatitis G virus.
GB virus (Appatitis G virus.
GBV-C/HGV group.
NIRL TAXID=54290;
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Pred. No. 3.6e+03;
2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 27; DB 2; Length 25;
Pred. No. 5.4e+03;
3; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kamo M., Kawakami T., Miyatake N., Tsugita A.;
Submitted (JUL-1994) to the PIR data bank.
PIR; PA0045; PA0045.

NON TER 1 1 1 1
SEQÜENCE 14 AA; 1546 MW; 0728ED7FB3BEBFBB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            25 25 25 25 AA; B67E75F5C0B77BF6 CRC64;
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01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Porin porl (Fragment).
Arabidopsis thaliana (Mouse-ear cress).
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Best Local Similarity 45.2
Best Local Similarity 45.2
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                  10 RRVDKDOWGPG 20
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 4 RKSERIORGPG 14
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Q7PE81
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"High prevalence of GB virus C strains genetically related to strains with Asian origin in Nicaraguan hemophiliacs.";
J. Med. Virol. 52:149-155(1997).
BMBL; U86i14; AABS8558.1; -.
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EMBL; U86:115; AAB58559.1; -.
                  Gaps
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MEDLINE=97323271; PubMed=9179760;
DOI=10.1002/(SICI)1096-9071(199706)52:2<149::AID-JMV5>3.3.CO;2-N;
GOnzalez-Perez M.A., Norder H., Bergetrom A., Lopez E., Visona K.A.,
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MEDLINE=97323271; PubMed=9179760;
DOI=10.1002/(SICI)1096-9071(199706)52:2<149::AID-JMV5>3.3.CO;2-N;
Gonzalez-Perez M.A., Norder H., Bergstrom A., Lopez E., Visona K.A.,
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Polyprotein (Fragment).
GB virus C/Hepatitis G virus.
Viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
GBV-C/HGV group.
NCBL_TaxID=54290;
                                                                                                                                                                                                                                                            GB virus C/Hepatítis G virus.
Viruses; ssRNA positive-strand viruses, no DNA stage; Flaviviridae;
GBV-C/HGV group.
NCBI_TaxID=54290;
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Pred. No. 5.4e+03;
3; Mismatches 3; Indels
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SEQUENCE 25 AA; 2800 MW; B67CADD5DBB77BF6 CRC64;
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Polyprotein (Fragment)
                  4;
 Pred. No. 5e+03;
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Similarity 46.2%; Pred. No. 5e+0 6; Conservative 3; Mismatches
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                                               4 RKSERIORGPGRA 16
                                                                     9 RRDKRRKRERGRA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5; Conservative
                                                                                                                                                                  PRELIMINARY;
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 Best Local Similarity
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NON TER
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Query Match
Beet Local Similarity 38.5.,
Trohes 5, Conservative
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nes 4; Conservative
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"Comparative study of protein profiles on pathogenic and nonpathogenic
Naegleria species by 2D-PAGE."
J. Bukaryot. Microbiol. 0:0-0:02004).
-!- MISCELLANEOUS: On the 2D-gel the determined pI of this unknown
protein is: 5.9, its MM is: 47.0 kDa.

Direct protein sequencing.

NON_TER
                                                                                                                                        Anopheles gambiae str. PEST.
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Anopheles.
NCBI_TaxID=180454;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Unknown protein NF041 from 2D-PAGE (Fragment).
Naegleria fowleri.
Eukaryota; Heterolobosea; Schizopyrenida; Vahlkampfiidae; Naegleria.
NCBI_TaxID=5763;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UNY-2003 (TrEMBLrel. 24, Last annotation update)
ALZHEIMER'S beta-amyloid precursor protein, kunitz-type protease inhibitor, neutrophil elastase inhibitor, Pl-VAL-APP-KD
                                                                                                                                                                                                                                                                                                                                   -i- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is preliminary data.

EMBL; AAAB0100434; EAA45843.1; -. SEQUENCE 14 AA; 1652 MW; 4A8A0A1AEC3F7FD3 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5; Indels
                                                                                                                                                                                                                                                                                              Anopheles Genome Sequencing Consortium;
Submitted (APR-2003) to the EMBL/GenBank/DDBJ databases.
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                                   01-WAR-2004 (TrEMBLrel. 26, Created)
01-WAR-2004 (TrEMBLrel. 26, Last sequence update)
01-WAR-2004 (TrEMBLrel. 26, Last annotation update)
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14 AA
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29-MAR-2004 (Rel. 43, Last seq
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PRELIMINARY;
                                                                                                                           Name=ENSANGG00000020916;
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                                                                                                     ENSANGP00000024647
                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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P83729;
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NP41_NREPO
DT 29-MAR,
DR 29-MAR,
DR 20-MAR,
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SEQUENCE FROM N.A.
STRAIN=DAEC18, DAEC19, DAEC2114, DAEC7, DAEC5, DAEC20,
DAEC162, DAECT2a, DAECT2b, DAEC11a, DAEC11b, DAEC318, DAEC719, EC7372,
ECOR37, DAEC9, ECOR50, IHIL128, C1845, and ECOR64;
PubMed=15014151; DOI=10.1093/molbev/mshil8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                        MEDLINE-92041969; PubMed-1939150; Sinha S., Knops J., Esch F., Moyer B.D., Oltersdorf T.; Sach F., Moyer B.D., Oltersdorf T.; Sach F., Moyer B.D., Oltersdorf T.; Moyer B.D., Oltersdon of the Alzheimer's beta-amyloid precursor protein (APP) Kunitz domain into a potent human neutrophil elastase inhibitor."; J. Biol. Chem. 266-211011-21013 (1991). GGO: GO: 0004867; F:serine-type endopeptidase inhibitor activity; NAS. GO: 00030162; P:regulation of proteolysis and peptidolysis; NAS.
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01-0CT-2003 (TrEMBLrel. 25, Last annotation update)
Gamma-glutam/ltranspeptidase heavy subunit (EC 2.3.2.2) (Fragment).
Bactllus subtilis.
Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus.
NCGI TaxID=1423;
Homo'sapiens (Human).
Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Enterobacteriaceae, Escherichia.
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Agric. Biol. Chem. 55:2971-2977 (1991).
GO, GO:0003840; F:gamma-glutamyltransferase activity; IEA.
SEQUENCE 17 AA: 1810 MW. 26198D040C958BEB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 26; DB 2; Length 17; Pred. No. 5.3e+03;
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Pred. No. 5.3e+03;
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Last sequence update)
Last annotation update)
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01-WAY-2000 (TrEMBLrel. 13, Last seq
01-OCT-2003 (TrEMBLrel. 25, Last ann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=92144110; PubMed=1371053;
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57.1%;
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38.5%;
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05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
25-OCT-2004 (TrEMBLrel. 28,
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EMBL, AYS04985; AAS46984.1; -. GO, GO:0016702; F:oxidoreductase activity, acting on single d. . .; II
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Ovis aries (Sheep).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
Caprinae; Ovis.
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Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
Pseudomonadaceae; Pseudomonas.
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Pred. No. 6.7e+03;
1; Mismatches 4; Indels
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Pred. No. 6.7e+03;
1; Mismatches 6; Indels
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TISSUE-Peripheral blood;
Elis S.A., Staines K.A.;
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ419229; CAD11671.1; -..
NON TER 1 1 1
SEQUENCE 21 AA; 2316 MW; BE2E264AACD38D6D CRC64;
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SEQÜENCE 21 AA; 2191 MW; 4200125E64F0F0DE CRC64;
                                                      01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DAP12 protein (Fragment).
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Catechol 2,3-dioxygenase (Fragment).
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01-0CT-2003 (TrEMBLrel. 25, Created)
01-0CT-2003 (TrEMBLrel. 25, Last sequence update)
21 AA.
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50.0%;
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54.5%;
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Best Local Similarity 54.5
Matches 6; Conservative
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  PRELIMINARY;
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S VPRGRGAAEVT 15
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Best Local Similarity
7; Conserve
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  Q95KS4
Q95KS4;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 25, Last annotation update)
Sulfite reductase 50 kDa alpha subunit (BC 1.8.99.3) (Fragment).
Desulfovibrio desulfuricana;
Desulfovibrionaceae; Desulfovibrio.
NCBI TAXID-876;
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"Molecular properties of the dissimilatory sulfite reductase from
Besulfovibrio desulfuricans (Essex) and comparison with the enzyme
from Desulfovibrio volgaris (Hildenborough).";
Bur. J. Blochem. 233:873-879(1995).
Bur. J. Blochem. 23490.
GO; GO:0018551; F:hydrogensulfite reductase activity; IEA.
SEQUENCE 20 AA; 2193 MW; F939B03B6B355135 CRC64;
                         Le Bouguenec C., Denamur E.;
"A Specific Genetic Background Is Required for Acquisition and
Expression of Virulence Factors in Escherichia coli.";
Mol. Biol. Evol. 21:1085-1094 (2004).
EMBL, AY525515; AAT00550.1; -.
EMBL, AY525517; AAT00556.1; -.
EMBL, AY525518; AAT00556.1; -.
EMBL, AY525519; AAT00556.1; -.
EMBL, AY525519; AAT00560.1; -.
EMBL, AY525519; AAT00560.1; -.
EMBL, AY525521; AAT00560.1; -.
EMBL, AY525521; AAT00560.1; -.
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     Escobar-Paramo P., Clermont O., Blanc-Potard A.B., Bui H.,
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Pred. No. 6.3e+03;
5; Mismatches 3; Indels
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MEDLINE=96085152; PubMed=8521853;
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EMBL, AY525524, AAT00566.11
EMBL, AY525246, AAT00570.11
EMBL, AY525226, AAT00570.11
EMBL, AY525226, AAT00574.11
EMBL, AY525229, AAT00576.11
EMBL, AY52529, AAT00576.11
EMBL, AY52531, AAT00580.11
EMBL, AY52531, AAT00581.11
EMBL, AY525331, AAT00581.11
EMBL, AY525331, AAT00581.11
EMBL, AY525334, AAT00588.11
EMBL, AY525531, AAT00588.11
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Local Similarity 33.3%;
les 4; Conservative :
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SEQUENCE FROM N.A.
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""" "The genome sequence of Schizosaccharomyces pombe.";

"" "The genome sequence of Schizosaccharomyces pombe.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
STRAIN=972h-;
MEDLINE=21848401; PubMed=11859360; DOI=10.1038/nature724;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 26; DB 2; Length 22;
Pred. No. 7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8; Indels
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Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
[2]
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                                                                                                          Schizosaccharomyces pombe (Fission yeast).
Bukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
Schizosaccharomycetales; Schizosaccharomycetaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
T17H7.9.
   01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nature 415:871-880(2002).
EMBL; AL158056; CAD99129.1; -.
GeneDB_SPombe; SPAC11H11.01; -.
                                          SPAC11H11.01 protein (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46.78;
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                                                                                                                                                                                                                             Schizosaccharomyces
                                                                    Name=SPAC11H11.01;
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                                                                                                                                                                                                                                                               NCBI_TaxID=4896;
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Plant Physiol. 133:1229-1239(2003).
EMBL; AX379780; AAQ74885.1; -
GO; GO:0047216; F:inositol 3-alpha-galactosyltransferase acti. .; IEA.
GO; GO:0016757; F:transferase activity, transferring glycosyl. .; IEA.
Glycosyltransferase; Transferase.
                                                                                                                                                                                                                                                                              Cheuk R., Shinn P., Brooks S., Buehler E., Chao Q., Johnson-Hopson C., Khan S., Kim C., Altafi H., Bei B., Chin C., Chiou J., Choi E., Conn L., Conway A., Gonzalez A., Hansen N., Howing B., Koo T., Lam B., Lie J., Lenz C., Li J., Liu A., Liu J., Liu S., Mukharsky N., Nguyen M., Palm C., Pham P., Sakano H., Schwartz J., Southwick A., Theologis A., Ecker J., Sakano H., Schwartz J., Southwick A., Theologis A., Ecker J., Sull M., Yu G., Davis R., Federspiel N., Theologis A., Ecker J., EmBL/GenBank/DDBJ databases.

EMBL, AC004135, Ab132934.1; --
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Momordica charantia (Bitter gourd) (Balsam pear).
Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnoliophyta, eudicotyledons; core eudicots; rosids;
eurosids I; Cucurbitales; Cucurbitaceae; Momordica.
Shinn P., Brooks S., Buehler E., Chao Q., Dunn P., Khan S., Kim C., Walker M., Altafi H., Araujo R., Conn L., Conway A., Gonzalez A., Hansen N., Huizar L., Kremenetskaia I., Enz C., Li J., Liu S., Luros S., Rowley D., Schwartz J., Toriumi M., Vysotskaia V., Yu, G, Davis R., Federapiel N., Theologis A., Ecker J., Submitted (MAY-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SECUENCE FROM N.A. MEDLINE-22975109; PubMed=14526110; DOI=10.1104/pp.103.027714; MEDLINE-22975109; PubMed=14526110; DOI=10.1104/pp.103.027714; Ayre B.G., Blair J.E., Turgeon R.; "Functional and phylogenetic analyses of a conserved regulatory "Functional and phylogenetic analyses of a conserved regulatory"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21.1%; Score 26; DB 2; Length 22; 62.5%; Pred. No. 7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Indels
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SEQUENCE 23 AA; 2444 MW; 62411699CAB81657 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          747DFB4CF6342ED6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JUL-2004 (TrEMBLrel. 27, Last sequence update) 05-JUL-2004 (TrEMBLrel. 27, Last annotation update) Galactinol synthase (EC 2.4.1.123) (Fragment).
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05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22 AA; 2642 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 62.5
nes 5; Conservative
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[1]
SEQUENCE FROM N.A.
MEDLINE-22975109; PubMed=14526110; DOI=10.1104/pp.103.027714;
MYPE B.G., Blair J.E., Turgeon R.;
Punctional and phylogenetic analyses of a conserved regulatory program in the phloem of minor veins.";
Plant Physiol. 133:1229-1239 (2003).
BMBL; AY379777; AAQ74879-13; -.
GO; GO:0047216; F:inositol 3-alpha-galactosyltransferase acti. . .; IEA.
GO; GO:0016757; F:transferase activity, transferase acti. . .; IEA.
GJ; GO:0016757; F:transferase.
NON TER
SEQÜENCE 24 AA; 2538 MW; E2BCOAE97330C06 CRC64;
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Galactinol synthase (EC 2.4.1.123) (Fragment).
Caterinol synthase (EC 2.4.1.123) (Fragment).
Caterilus lanatus (Watermelon) (Citrullus vulgaris).
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
euroalds I; Cucurbitales; Cucurbitaceae; Citrullus.
NCBI_TaxID=3654;
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              DE REPRESENTATION OF STREET OF STREE
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0; Gaps

Search completed: May 16, 2005, 13:00:27 Job time : 138.231 secs

13 PGRAFVT 19 | ||:|| 17 PKRAYVT 23

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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

- protein search, using sw model OM protein Run on:

May 16, 2005, 12:53:32 ; Search time 28.9231 Seconds (without alignments) 79.839 Million cell updates/sec

US-08-869-386-3

123 1 NNTRKSERIQRGPGRAFVTIGKIG 24 Perfect score: Sequence:

Gapop 10.0 , Gapext 0.5 BLOSUM62 Scoring table:

283416 seqs, 96216763 residues Searched:

4989 Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 25 Maximum DB & Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 79:\*
1: pir1:\*
3: pir2:\*
1: pir3:\* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	immunodeficiency	hydrogensulfite re	MSEL-neurophysin	Plasmepsin II - m	T-cell receptor al	por 1B	porin porl - Arabi	dissimilatory sulf	protein T17H7.9	protein P7 - curle	exo-alpha-sialidas	translation initia	heat shock protein	ribosomal protein	dihydrolipoamide S	T-cell receptor be	seed protein ws-2	Ig kappa-1 chain	T-cell receptor be	lamin B receptor		probable angiotens		hypothetical prote		36K protein 3124	diaminopropionate	m	cytochrome-c oxida
	865399	S21197	B60422	S48654	803505	PA0109	PA0045	S63490	H86433	D28027	S11286	JN0264	A60807	S51071	A37823	PT0515	G61497	C53275	PH0915	S28405	3142	B48395	P41476	C42856	D41575	PS0453	Ã.	T01780	865388
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integration host f fibrinopeptide A - myosin heavy chain heterocyst-inhibit translation elonga TcR delta chain V- tet leader peptide cytochrome-c oxida ribosomal protein nucleoside-diphosp protoporphyrinogen T cell receptor al T-cell receptor al glucose 1-dehydrog cAMP-regulated pho	a (TIAGGG) n-binaing
PN0629 H29501 H29501 AF2033 S39153 S39153 F149037 EF8TT S77991 S65623 S65629 S65629 S66613 S66613 S66613 S66613 S66613	B48138
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<b>0 11 12 14 14 14 14 14 14 14 14 14 14 14 14 14 </b>	45

## ALIGNMENTS

Cispectes: Homo sapiens (man)
Cispectes: Homo sapiens (man)
Cispectes: Homo sapiens (man)
Cipate: 28-Oct-1996 #sequence\_revision 13-Mar-1997 #text\_change 17-Mar-1999
Cipacession: S65399
Airitle: Treel membrane-associated serine protease, tryptase TL(2), binds human immunode man immunodeficiency virus type 1 inhibit cleavage of gpl20.
Airitle: Treel member: S65399, MUID:96203909, PMID:8620895
Airitle arguerianiary
A immunodeficiency virus type 1, HIV-1 gp120 - human (fragments)

Gaps ; 0 Score 64; DB 2; Length 20; Pred. No. 0.0017; 1; Mismatches 0; Indels 52.0%; 92.3%; Query Match 52.0 Best Local Similarity 92.3 Matches 12; Conservative

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11 RGPGRAFVTIGKI 23 1 RGPGRAFVTIGRI 13 ð a

RESULT 2 S21197

hydrogensulfite reductase (BC 1.8.99.3) alpha chain - Desulfovibrio vulgaris (fragment)
Nylternate names: bisulfite reductase; desulfofuscidin; desulforubidin; desulfoviridin;
C;Species: Desulfovibrio vulgaris
C;Date: 19-Mar-1997 #sequence\_revision 11-Jun-1999 #text\_change 09-Jul-2004
C;Accession: S21197
R;Pierik, A.J.; Duyvis, M.G.; van Helvoort, J.M.L.M.; Wolbert, R.B.G.; Hagen, W.R.
Bur. J. Biochem. 205, 111-115, 1992
A;Title: The third subunit of desulfoviridin-type dissimilatory sulfite reductases.
A;Reference number: S21197; MUD:92209491; PMID:1555572
A;Accession: S21197
A;Molecule type: protein
A;Residues: 1-25 \*PIB>
A;Cross-references: UNIPROT:P45574
A;Experimental source: strain Hildenborough
C;Genetics:
C;Genetics:
C;Complex: heterohexamer; two alpha, two beta and two gamma chains
C;Complex: heterohexamer; two alpha, two beta and two gamma chains

A; Description: catalyzes the six-electron reduction of sulfite to sulfide A; Pathway: the terminal oxidase in the sulfate-reduction pathway C; Keywords: heterohexamer; oxidoreductase

Length 25; 26.8%; Score 33; DB 2; I 37.5%; Pred. No. 1.7e+02; Query Match Best Local Similarity

Matches

RESULT 3

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dissimilatory sulfite reductase alpha chain, soluble - Desulfovibrio desulfuricans (fragm C;Species: Desulfovibrio desulfuricans (5) Species: Desulfovibrio desulfuricans (5) Species: 28-0ct-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004 C;Accession: S63490 R;Steuber, J.; Arendsen, A.F.; Hagen, W.R.; Kroneck, P.M.H. Bur. J. Biochem. 233, 873-879, 1995 A;Tile: Molecular properties of the dissimilatory sulfite reductase from Desulfovibrio characterism number: S63489; MuID:96085152; PMID:8521853
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 30-unn-1992 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
C;Datesion: B40045
R;Kamo, M.; Kawakami, T.; Miyatake, N.; Tsugita, A.
submitted to JIPID, July 1994
A;Description: Separation and characterization of Arabidopsis proteins by two-dimensional
A;Reference number: PA0001
                                                                                                                                                                                                                                                                         porin por 1B - Arabidopsis thaliana (fragment)
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 07-Apr-1995 #sequence_revision 26-May-1995 #text_change 09-Jul-2004
C;Accession: PAOLDS
R;Kamo, M; Kawakami, T.; Tsugita, A.
Bibmitted to JIPLD, March 1995
A;Reference number: PAO109
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Pred. No. 1.1e+03;
2; Mismatches 3;
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Pred. No. 1.1e+03;
                    22.0%; Score 27; DB 2; I
29.4%; Pred. No. 1.2e+03;
iive 4; Mismatches 8;
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A; Molecule type: protein
A; Residues: 1-14 < KAAN>
A; Cross-references: UNIPROT: Q8LA84; UNIPROT: Q42292
A; Experimental source: root
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A, Experimental source: root
                                                                                                                      2 NTRKSERIQRGPGRAFV 18
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Best Local Similarity 50.0
Matches 6; Conservative
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Best Local Similarity
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A;Molecule type: protein
A;Residues: 1-20 <STE>
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R;Hill, J.; Tyas, L.; Phylip, L.H.; Kay, J.; Dunn, B.M.; Berry, C.
BESS Lett. 352, 155-158, 1994
A;Title: High level expression and characterisation of Plasmepsin II, an aspartic protein A;Reference number: S48654; MUID:95010698; PMID:7925966
                                                                                                                                                                                                                                          NiAlternate names: vasopressin-associated neurophysin
C;Species: Xenopus laevis (African clawed frog)
C;Date: 12-Peb-1993 #sequence_revision 12-Peb-1993 #text_change 17-Mar-1999
C;Accession: B60422
R;Chauvet, J; Michel, G; Rouille, Y; Chauvet, M.T.; Acher, R.
Neuropeptides 15, 123-127, 1990
A;Title: Identification of two types of neurophysins in Xenopus laevis neurointermediate
A;Reference number: A60422; MUID:91067001; PMID:2250763
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C;Species: Mus*musculus (house mouse)
C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change 30-May-1997
C;Accession: 803505
R;Winoto, A.; Mjolsness, S.; Hood, L.
Nature 316, 832-836, 1985
A;Title: Genomic organization of the genes encoding mouse T-cell receptor alpha-chain. A;Reference number: 803503; MUID:85296332; PMID:2993908
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C;Species: Plasmodium falciparum
C;Date: 15-Jul-1995 #sequence_revision 19-Apr-1996 #text_change 09-Jun-2000
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Pred. No. 1.2e+03;
1; Mismatches 5; Indels
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A,Residues: 1-20 <WIN>
A,Cross-references: EMBL:X02859
A,Note: this sequence was determined from the germline gene
C,Keywords: T-cell receptor
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Pred. No. 1e+03;
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MSEL-neurophysin - African clawed frog (fragment)
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A;Residues: 1-24 <CHA>
C;Superfamily: oxytocin-neurophysin
C;Keywords: pitultary
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5 TPKLDQLESGPWXSFV 20
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Local Similarity 38.5%;
nes 5; Conservative
                                                       3 TRKSERIORGPGRAFV
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Best Local Similarity 50.0
Matches 6; Conservative
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            Conservative
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A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-20 <HIL>
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A;Molecule type: protein
A;Residues: 1-25 <DEN>
C;Comment: This protein associates with steroid hormone receptors and with the Rous sarcc
C;Superfamily: heat shock protein 90
C;Keywords: phosphoprotein
                            C;Species: influenza A virus
C;Species: influenza A virus
C;Date: 31-Dec-1990 #sequence_revision 31-Dec-1990 #text_change 22-Jun-1999
C;Date: 31286
R;Robertson, J.S.
Nucleic Acids Res. 6, 3745-3757, 1979
A;Title: 5' and 3' terminal nucleotide sequences of the RNA genome segments of influenza A;Reference number: S1286; MUID:80034428; PMID:493121
A;Reference number: S1286
A;Molecule type: genomic RNA
A;Residues: 1-12 <ROB>
A;Cross-references: EMBL:J02114; NID:g324483; PIDN:AAA43398.1; PID:g324486
C;Genetics:
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NyAlternate names: eIP2 gamma chain
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 09-Jul-2004
C;Accession: JN0264 #sequence_revision 09-Oct-1992 #text_change 09-Jul-2004
C;Accession: JN0264 #sequence_revision 09-Oct-1992 #text_change 09-Jul-2004
C;Accession: JN0264 #sequence in the gamma subunit of pig liver initiation factor 2.
A;Reference number: JN0264; MUID:92282179; PMID:1368212
A;Accession: JN0264
A;Accession: JN0264
A;Accession: JND264
A;Accesi
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C;Species: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 17-Mar-1999
C;Accession: A60807
R;Denis, M.
Anal. Biochem. 173, 405-411, 1988
A;Title: Two-step purification and N-terminal amino acid sequence analysis of A;Reference number: A60807; MUID:89048319; PMID:3189918
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Pred. No. 2.1e+03;
2; Mismatches 3; Indele
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Pred. No. 3.2e+03;
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Pred. No. 1.6e+03;
2; Mismatches 3
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C;Superfamily: influenza virus exo-alpha-sialidase
C;Keywords: glycosidase; hydrolase
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N, Alternate names: neuraminidase
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Best Local Similarity 44.4
Matches 4; Conservative
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Proc. Natl. Acad. Sci. U.S.A. 84, 4806-4810, 1987
A;Title: Alterations in the phenotype of plant cells studied by NH2-terminal amino acid-A;Reference number: A94167
A;Accession: D28027
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein T17H7.9 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar_2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C,Species: Nicotiana plumbaginifolia (curled-leaved tobacco)
C,Date: 19-May-1989 #sequence_revision 19-May-1989 #text_change 09-Jul-2004
C,Accession: D28027
                                                                                                                                                       Gaps
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                                    Score 26; DB 2; Length 2u;
Pred. No. 1.88+03;
"..marrhes 3; Indels
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Pred. No. 1.3e+03;
1; Mismatches 1;
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Pred. No. 2e+03;
1; Mismatches
                                                                                                                                                       5; Mismatches
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A,Residues: 1-10 <BAU>
A,Cross-references: UNIPROT:Q7M1V8
A; Cross-references: UNIPROT: Q9R4H4
                                                                 21.1%;
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62.5%;
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Best Local Similarity 71.4
Matches 5; Conservative
                                                                                                              Local Similarity 33.3
hes 4; Conservative
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                                                                                                                                                                                                                                7 ERIQRGPGRAFV 18
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NKKKKSER 15
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Best Local Similarity
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A;Residues: 1-22 <STO>
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                                                                             Query Match
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T.cell receptor beta chain V-D-J region (isolate 1) - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 09-0ct-1992 #sequence_revision 09-0ct-1992 #text_change 30-May-1997
C;Accession: PH0915
R;Gold, D.P.; Offiner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
J. Exp. Med. 174, 1467-1457, 1991
A;Fitle: Analysis of T cell receptor beta chains in Lewis rats with experimental allergit A;Reference number: PH0891; MUID:92078857; PMID:1836012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Residues: 1-13 <AYA>
A;Note: sequence extracted from NCBI backbone (NCBIN:56069, NCBIP:56164)
C;Comment: This J3 segment may not be functional because of substitutions in the 7 mer ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Title: Microsequence analysis of winged bean seed proteins electroblotted from two-dime A;Reference number: A61491; MUID:89351606; PMID:2765119
A;Accession: G61497
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C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: (2-May-1994 #sequence_revision 18-Nov-1994 #text_change 16-Aug-1996
                                                                                                                                                                                                                                                                                                           seed protein ws-23 - winged bean (fragment)
C;Species: Psophocarpus tetragonolobus (winged bean)
C;Date: 07-Oct.1994 #sequence_revision 07-Oct.1994 #text_change 07-Oct-1994
C;Accession: G61497
K;Hirano, H.
J. Protein Chem. 8, 115-130, 1989
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R;Ayadi, H.; Marche, P.N.; Cazenave, P.A.
RiAyadi, H.; Marche, P.N.; Cazenave, P.A.
Rimunogenetics 4, 201-207, 1991
A;Title: Evolution of the rabbit immunoglobulin kappa chain genes.
A;Reference number: A53275; WUID:91372868; PMID:1909995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19.5%; Score 24; DB 2; Length 13; 100.0%; Pred. No. 2.5e+03;
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                                              Score 24; DB 2; I
Pred. No. 2.8e+05;
1; Mismatches 0;
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Pred. No. 2.1e+03;
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50.0%;
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A;Molecule type: protein
A;Residues: 1-11 <HIR>
C;Keywords: glycoprotein; seed
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                                              Query Match
Best Local Similarity 80.0
Matches 4; Conservative
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Matches 5; Conservative
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C; Keywords: T-cell receptor
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Matches 4; Conser
                                                                                                                                          12 GPGRA 16
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A;Molecule type: DNA
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PH0915
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C)Accession: 851071
Shysiboli, P.; Herfurth, E.; Choli, T.
Bur. J. Blochem. 226, 169-177, 1994
A;Title: Purification and characterization of the 30S ribosomal proteins from the bacter A;Reference number: 851053; MUD:95045586; PMID:7957245
A;Accession: S51051
A;Molecule type: protein
A;Residues: 1-25 cTS1>
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A;Title: Changes in the core of the mammalian-pyruvate dehydrogenase complex upon select A;Reference number: A37823; MJID:90354445; PMID:2167319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R.Feeney, A.J. 4, 115-124, 1991 A.J. Ecall receptor beta chains have few N regions. A.File: Junctional sequences of fetal T call receptor beta chains have few N regions. A.Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0515
                                                                                                                                                                                                                                                     C;Species: Thermus aquaticus
C;Date: 15-Jul-1995 #sequence_revision 01-Nov-1996 #text_change 01-Nov-1996
C;Accession: S51071
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     IndelB
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Pred. No. 3.2e+03;
2; Mismatches 7;
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66.7%; Pred. No. 2.6e+03;
iive 1; Mismatches 1;
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                                                                                                                                                                                                                            ribosomal protein S20 - Thermus aquaticus (fragment)
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     Mismatches
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A,Residues: 1-17 «RAH»
A,Cross-references: UNIPROT: G7M2M8
C;Keywords: acyltransferase; coenzyme A
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ilarity 35.7%;
Conservative ;
                                                     7 ERIORG--PGRAFVT 19
                                                                                                   EEVQKGEEPVETFAT 16
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A;Steuus: translation not shown
A;Molecule type: mRNA
A;Residues: 1-7 <FEE>
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       Conservative
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Matches 5; Conserv
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Best Local Similarity
Matches 6; Conserv
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A; Status: preliminary
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C;Date: 21-Jan-1994 #sequence_revision 23-Mar-1995 #text_change 09-Jul-2004
C;Accession: B48395
R;Maruyama, E.; Iwamatsu, A.; Takashima, S.
Biocohem. Mol. Biol. Int. 29, 579-586, 1995
A;Title: Purification and amino acid microsequencing of alkaline phosphodiesterase I from A;Title: Purification and amino acid microsequencing of A;Accession: B48395
A;Status: preliminary
A;Status: preliminary
A;Nolecule type: protein
A;Residues: 1-22 <MAR>
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CjSpecies: Homo sapiens (man)
CjSpecies: Homo sapiens (man)
CjAccesion: C42856
R;Liu, Z.; Diaz, L.A.; Haas, A.L.; Giudice, G.J.
J. Biol. (Chem. 267, 15829-15835, 1992
A;Title: cDNA cloning of a novel human ubiquitin carrier protein. An antigenic domain spethis human epidermal transcript.
A;Reference number: A42856; MUID:92348449; PMID:1379239
A;Accession: C42856
A;Actus: preliminary
A;Molecule type: mRNA
A;Residues: 1-22 cLIU>
A;Reperimental source: keratinocyte
A;Reperimental source: keratinocyte
A;Note: sequence extracted from NCBI backbone (NCBIN:109895, NCBIP:109899)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probable antigen 6 - Mycobacterium leprae (fragment)
C;Species: Mycobacterium leprae
C;Date: 10-Apr-1992 #sequence_revision 10-Apr-1992 #text_change 18-Jun-1993
C;Accession: F41476
R;Hartskeerl, R.A.; van Rens, R.M.; Stabel, L.F.E.M.; de Wit, M.Y.L.; Klatser, P.R.
Infect. Immun. 58, 2821-2827, 1990
A;Title: Selection and characterization of recombinant clones that produce Mycobacterium A;Reference number: A41476; WUID:90354041; PMID:1696931
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A;Residues: 1-22 <HAR>
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Best Local Similarity 37.5%; Pred. No. 4e+03;
Matches 6; Conservative 2; Mismatches
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Pred. No. 4e+03;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:Q9TRH0
A;Experimental Source: Kidney
A;Note: sequence extracted from NCBI backbone
C;Superfamily: mammalian peptidyl-dipeptidase A
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Best Local Similarity 100..
Laga 4; Conservative
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Best Local Similarity 50.0
Matches 5; Conservative
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R;Simos, G.; Georgatos, S.D.
EMBO J. 11, 4027-4036, 1992
A;Title: hiner nuclear membrane protein p58 associates in vivo with a p58 kinase and A;Reference number: S28405; MUID:93010998; PMID:1327755
                  A;Accession: PH0915
A;Molecule type: mRNA
A;Residues: 1-14 <GOL>
C;Residues: 1-14 <GOL>
C;Residue B as Glu and GAG for residue B codon GAG for residue B c
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C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 07-Feb-1997
C;Accession: S31427.
A;Nedellec, P.; Turbide, C.; Barnett, T.R.; Beauchemin, N.
submitted to the EMBL Data Library, July 1992
A;Description: Characterization of the human biliary glycoprotein regulatory region.
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NiAlternate names: inner nuclear membrane protein p58
CiSpecies: Meleagris gallopavo (common turkey)
CiSpecies: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
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A, Molecule type: protein
A, Molecule type: protein
A, Residues: 1-20 «SIM»
A, Cross-references: UNIPROT: Q7LZ11
C, Keywords: DNA binding; nucleus; receptor; transmembrane protein
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                                                                                                                                                                                                                                                                     Length 14;
                                                                                                                                                                                                                                                                 19.5%; Score 24; DB 2; Length 14; 50.0%; Pred. No. 2.6e+03; ative 2; Mismatches 2; Indels
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C,Species: Bos primigenius taurus (cattle)
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Best Local Similarity 80.0
Matches 4; Conservative
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Best Local Similarity 50.0
Matches 4; Conservative
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A, Accession: S31427
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A;Residues: 1-21 <NED>
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cytochrome-c oxidase (EC 1.9.3.1) chain VII c, hepatic - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 28-Oct-1996 Hesquence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Date: 28-Oct-1996 Hesquence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: S65388; S6538
R;Schaegger, H.; Noack, H.; Halangk, W.; Brandt, U.; von Jagow, G.
Bur. J. Blochem. 230, 235-241, 1995
A;Title: Cytochrome-c oxidase in develOping rat heart. Enzymic properties and amino-term;
A;Reference number: S65372; MUD:95324529; PMID:7601105
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C;Species: Pseudomonas aeruginosa
C;Date: 05-Aug-1994 #sequence_revision 05-Aug-1994 #text_change 09-Jul-2004
C;Date: 05-Aug-1994 #sequence_revision 05-Aug-1994
R;Date: 05-Aug-1994 #sequence_revision 1: // Vignais, P.M.
Biochem. Biophys. Res. Commun. 196, 416-421, 1993
A;Title: Pseudomonas aeruginosa contains an IHF-like protein that binds to the algD promt A;Reference number: PN0629; MUID:94030028; PMID:8216322
A;Accession: PN0629
A;Molecule type: protein
                                                                                                                                                                                                                                                                     R;Repaske, R.; O'Neill, R.R.; Steele, P.E.; Martin, M.A.
Proc. Natl. Acad. Sci. U.S.A. 80, 678-682, 1983
A;Title: Characterization and partial nucleotide sequence of endogenous type C retrovirus
A;Reference number: Z14423; MUID:83143394; PMID:6298769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    integration host factor-like protein beta chain - Pseudomonas aeruginosa (fragment)
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                                                                                                                                                       probable gag polymerase pseudogene - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 13-Aug-1999 #sequence_revision 13-Aug-1999 #text_change 20-Oct-2000
C;Accession: T01780
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A;Molecule type: protein
A;Residues: 1-10 <SCH>
A;Cross-references: UNIPROT:P80432
A;Actession: 865389
A;Molecule type: protein
A;Residues: preliminary
A;Residues: 1-10 <SC2>
C;Superfamily: cytochrome-c oxidase chain VIIc
C;Keywords: oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Cross-references: EMBL: J00274; NID: 9182154
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10 NTR---RKKKGTGAA 21
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Best Local Similarity 33.33
Matches 6; Conservative
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Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Keywords: pseudogene
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PN0629
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                               셤
                                                                                               bombinin-like peptide 4 - Bombina orientalis
C;Species: Bombina orientalis
C;Species: Bombina orientalis
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 09-Jul-2004
C;Accession: D41575
C;Accession: D41575
J. Biol. Chem. 266, 23103-23111, 1991
J. Biol. Chem. 266, 23103-23111, 1991
A;Fitle: Bombinin-like peptides with antimicrobial activity from skin secretions of the A;Reference number: A41575; MUID:92078177; PMID:1744108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N;Alternate names: diaminopropionatase
C;Species: Salmonella typhimurium
C;Date: 28-Aug-1989 #sequence_revision 28-Aug-1989 #text_change 18-Jun-1993
C;Accession: A28524
R;Nagasaa, T: ? Tanizawa, K.; Satoda, T.; Yamada, H.
J. Biol. Chem. 263, 958-964, 1988
A;Title: Diaminopropionate ammonia-lyase from Salmonella typhimurium. Purification and
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C;Species: Oryza sativa (rice)
C;Species: Oryza sativa (rice)
C;Accession: PS0453
R;Tsugita, A.
submitted to JIPID, April 1993
A;Reference number: PS0206
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46.7%; Pred. No. 4.8e+03;
tive 2; Mismatches 3; Indels
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Pred. No. 4.5e+03;
2; Mismatches 4; Indels
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19.1%; Score 23.5; DB 2;
Best Local Similarity 50.0%; Pred. No. 2.9e+03;
Matches 6; Conservative 1; Mismatches 4;
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A,Residues: 1-13 «TSU»
A,Experimental source: leaf, chlorophyll, stem
A,Note: molecular weight 36K, pI 6.1
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C; Superfamily: bombinin H precursor
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Similarity 45.5%;
5; Conservative 2
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A28524
diaminopropionate ammonia-lyase
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Best Local Similarity 46.7
Matches 7; Conservative
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GIGAAILSAGK 11
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A,Molecule type: protein
A,Residues: 1-22 <NAG>
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Best Local Similarity
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RESULT 26 PS0453

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A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium Anak A;Reference number: AB1807; MUID:21595285; PMID:11759840
A;Accession: AF2093
A;Status: preliminary
A;Status: preliminary
A;Residues: DNA
A;Residues: 1-17 < KUR>
A;Residues: 1-17 < KUR>
A;Cross-references: UNIPROT:052748; GB:BA000019; PIDN:BAB74000.1; PID:g17131393; GSPDB:G}
C;Genetics:
A;Gene: patS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   translation elongation factor EF-Tu, chloroplast - common tobacco (fragment)
C;Species: Nicotiana tabacum (common tobacco)
C;Species: Nicotiana tabacum (common tobacco)
C;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 05-Dec-1997
C;Accession: S39153
R;Murayama, Y.; Matsubayashi, T.; Sugita, M.; Sugiura, M.
Plant Mol. Biol. 22, 767-774, 1993
A;Title: Purification of chloroplast elongation factor Tu and cDNA analysis in tobacco: t
A;Reference number: S36183; MUID:93363910; PMID:8358028
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REZQUETRA, A.; Wilde, D.B.; McConnell, T.J.; Sturmhofel, K.; Valas, R.B.; Shevach, E.M.;
Rur. J. Immunol. 22, 491-498, 1992
A.Title: Mouse autoreactive gamma/delta T cells. II. Molecular characterization of the T A;Reference number: A49037; MUD:92164730; PMID:1311262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-18 <MUR>
C;Superfamily: translation elongation factor Tu; translation elongation factor Tu homolog
C;Keywords: chloroplast; GTP binding; protein biosynthesis
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TCR delta chain V-D-J region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 05-Nov-1999
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A;Experimental source: dendritic epidermal T-cell lines
A;Note: sequence extracted from NCBI backbone (NCBIN:90660, NCBIP:90671)
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Pred. No. 4.5e+03;
1; Mismatches 1; Indels
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Pred. No. 4.8e+03;
0; Mismatches 2;
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Matches 6; Conservative
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Best Local Similarity
Matches 4; Conserv
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12 ERGSGR 17
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C;Date: 21-Nov-1987 #sequence_revision 21-Nov-1987 #text_change 09-Jul-2004
C;Accession: H29501
R;Blombacck, B.; Blombacck, M.; Hann, C.
unpublished results, cited by Blombacck, B., and Blombacck, M., in Chemotaxonomy and Sex A;Reference number: A29501
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A;Molecule type: protein
A;Residues: 1-16 <BLO>
A;Cross-references: UNIPROT:Q7M316
C;Superfamily: fibrinogen beta chain; fibrinogen beta/gamma homology; fibrinogen disulfi
     A;Residues: 1-15 <TOU>
A;Cross-references: UNIPROT:Q9R533
C;Comment: This protein forms a stable complex with the algD promoter in vitro, indicati
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R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, A.; Iriguchi
Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada, M.; Yasuda, M.; Tabata, S
DNA Res. 8, 205-213, 2001
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C;Species: Gallus gallus (chicken)
C;Species: Gallus gallus (chicken)
C;Date: 04-Sep-1997 #sequence_revision 07-Nov-1997 #text_change 09-Jul-2004
C;Accession: IS1203
R;Kelley, C.A.; Takahashi, M.; Yu, J.H.; Adelstein, R.S.
J. Biol. Chem. 268, 12848-12854, 1993
A;Telle: An insert of seven amino acids confers functional differences between smooth in A;Reference number: IS1203; MUID:93286132; PMID:8509418
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C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC 7120
C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 09-Jul-2004
C;Accession: AF2093
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A;Molecule type: mRNA
A;Residues: 1-17 <KEL>
                                                                                                          Score 23; DB
Pred. No. 4e+0
1; Mismatches
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Best Local Similarity 71.4%;
Matches 5; Conservative 1
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Best Local Similarity 30.8%;
Matches 4; Congeresting
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                                                                                                                                                                                                                                                                                                                                                                                              - gray seal
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KSELIER 9
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Best Local Similarity
Matches 4; Conserv
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ribosomal protein S19 - evening primrose mitochondrion (fragment)
c;bosomal protein S19 - evening primrose)
c;bate: 10-Dec-1994 #sequence_revision 12-May-1995 #text_change 12-Jun-2003
c;bate: 10-Dec-1994 #sequence_revision 12-May-1995 #text_change 12-Jun-2003
c;Accession: S43765; S30542
R;Bock, H:, Brennicke, A.; Schuster, W.
Plant Mol. Biol. 24, 811-818, 1994
A;Title: Rps3 and rpl16 genes do not overlap in Oenothera mitochondria: GTG as a potential A;Reference number: S43765; MUID: 94250844; PMID: 8193306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Avena sativa (oat)
C;Species: Avena sativa (oat)
C;Date: 07-May-1995 #sequence_revision 24-Oct-1997 #text_change 09-Jul-2004
C;Accession: 847563
R;Sommer, D.; Song, P.S.
Biochim. Biophys. Acta 1222, 464-470, 1994
A;Title: A plant nucleoside diphosphate kinase homologous to the human Nm23 gene product: A;Reference number: 847563; MUD:94312444; PMID:8038216
                                                               A;Genome: nuclear
C;Function:
A;Pathway: oxidative phosphorylation; respiratory chain
C;Keywords: electron transfer; membrane-associated complex; mitochondrial inner membrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Accession: S41765
A;Molecule type: DNA
A;Residues: 1-23 BOC>
A;Cross-references: BMBL:X69140; NID:g13188; PIDN:CAA48893.1; PID:g13189
A;Note: the source is designated as Oenothera berteriana
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C,Superfamily: nucleoside diphosphate kinase
C,Keywords: phosphotransferase; pyrimidine nucleotide biosynthesis
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Pred. No. 6.3e+03;
2; Mismatches 2; Indels
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Similarity 40.0%; Pred. No. 5.3e+03;
4; Conservative 2; Mismatches 4; Indels
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A;Genome: mitochondrion
C;Superfamily: ribosomal protein S19/S15
C;Keywords: mitochondrion; protein biosynthesis; ribosome
A, Experimental source: heart, liver C, Genetics:
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50.0%;
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Best Local Similarity 41.7
Matches 5, Conservative
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Best Local Similarity 50.0
Matches 4; Conservative
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A, Molecule type: protein
A, Residues: 1-24 < SOM>
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Best Local Similarity
Matches 4; Conserv
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R; Kunst, F; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Berter
R; Kunst, F.; Ogasawara, N.; Moszer, I.; Albertini, B.; Capuano, V.; Carter, N.M.; Chc
C; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chc
A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A; Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Galler
iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, M.; Kurita, K.; Lapidus, A.; Lullo, M.F.
Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois,
Koetter, P.; Koningstein, G.; Krogh, S.; Musel, M.; Muthors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maucel,
Y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon,
A; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Seror
akeuchi, M.; Tamakoshi, A.; Tamaka, T.; Terpstray, P.; Tognoni, A.; Tosato, V.; Uchiyama,
T.; Winters, P.; Winters, P.; Winters, P.; Yamamoto, H.; Yamane, K.; Yasumoto, M.; Anthors: Voblida, M.; A.; Accession: H69721
A; Accession: H69721
A; Statumoto, R.; Andelic caid sequence not shown; translation not shown
                         tet leader peptide - Bacillus subtilis
N;Alternate names: tetracycline resistance leader peptide tetL
C;Species: Bacillus subtilis
C;Species: Bacillus subtilis
C;Date: 30-Jun-1991 #sequence revision 30-Jun-1991 #text_change 09-Jul-2004
C;Accession: 804802; 517282; H69721
R;Sakaguchi, R.; Amano, H.; Shishido, K.
Biochim. Biophys. Acta 950, 441-444, 1988
A;Title: Nucleotide sequence hoology of the tetracycline-resistance determinant natural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A,Cross-references: UNIPROT:P23053; EMBL:X08034; NID:g40207; PIDN:CAA30826.1; PID:g40208 R;Amano, H.; Sakaguchi, R.; Shishido, K. FEMS Microbiol. Lett. 79, 5-8, 1991 A;Title: An insertion of Escherichia coli transposable element ISIK into the site immedi
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A;Experimental source: strain 168
Genetics:
A;Gene: tetL
C;Superfamily: tet leader peptide
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877991
cytochrome-c oxidase (EC 1.9.3.1) chain VIII.1 - bigeye tuna (fragment)
C;Species: Thunnus obesus (bigeye tuna)
C;Bate: 17-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 09-Jul-2004
C;Accession: S77991
C;Accession: S77991
S;Arnold, S.; Lee, J.; Kim, M.; Song, E.; Linder, D.; Lottspeich, F.; Kadenbach, Submitted to the Protein Sequence Database, June 1997
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                                                                                                                                                                                                                                                                                                                                                            A;Reference number: JT0427; MUID:89000797; PMID:2844262 A;Accession: S04802
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A,Accession: S77991
A,Molecule type: PANO-en
A,Residues: 1-20 <ARN-A,Residues: A-ARN-A,COSS-references: UNIPROT:P80983
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A; Residues: 1-20 < KUN>
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A; Residues: 1-20 <AMA>
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Jucose 1-dehydrogenase (EC 1.1.1.47) - Haloferax mediterranei (fragment)
C;Species: Haloferax mediterranei
C;Species: Haloferax mediterranei
C;Date: 14-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: S66113
R;Bonete, M.J.; Pirt. C.; LLorca, F.I.; Camacho, M.L.
R;Bonete, M.J.; Pirt. C.; LLorca, F.I.; Camacho, M.L.
A;Fitle: Glucose dehydrogenase from the halophilic Archaeon Haloferax mediterranei: enzym
A;Reference number: S66213; MUID:96198607; PMID:8925901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAMP-regulated phosphoprotein, 21K - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Accession: A33361
R;Hemmings Jr., H.C.; Girault, J.A.; Williams, K.R.; LoPresti, M.B.; Greengard, P. J. Biol. Chem. 264, 7726, 7731
J. Biol. Chem. 264, 7736, 7739
A;Title: ARPP-21, a cyclic AMP-regulated phosphoprotein (M-r=21,000) enriched in dopamine netic studies of its phosphorylation in vitro.
A;Reference number: A33361; MUID:89214228; PMID:2540203
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C;Species: Homo sapiens (man)
C;Date: 16-Feb-1994 #sequence_revision 18-Nov-1994 #text_change 16-Aug-2004
C;Accession: B48138
R;Ishikawa, F.; Matunis, M.J.; Dreyfuss, G.; Cech, T.R.
Mol. Cell. Biol. 13, 4301-4310, 1993
A;Title: Nuclear proteins that bind the pre-mRNA 3' splice site sequence r(UUAG/G) and t)
A;Reference number: A48138; MUID:93309464; PMID:8321232
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N,Alternate names: type B heterogeneous nuclear ribonucleoprotein homolog
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Pred. No. 6.5e+03;
3; Mismatches 6; Indels
Indels
9
Mismatches
  4
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A;Residues: 1.17 *BON>
A;Cross-references: UNIRBOT:0977U7
C;Keywords: oxidoreductase
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Best Local Similarity 35.7%;
Matches 5; Conservative
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Best Local Similarity 26.77
Matches 4; Conservative
  4; Conservative
                                                                                       2 DKVIFGPGTSLSVI
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A; Status: preliminary
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                                                                                                                                   C;Accession: S65629
R;Taketani, S.; Yoshinaga, T.; Furukawa, T.; Kohno, H.; Tokunaga, R.; Nishimura, K.; Inc
Biochem. 230, 760-765, 1995
A;Title: Induction of terminal enzymes for heme biosynthesis during differentiation of m
A;Reference number: 865629; MUID:95331315; PMID:7607249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Accession: H41299
R;Uematsu, Y.; Wege, H.; Straus, A.; Ott, M.; Bannwarth, W.; Lanchbury, J.; Panayi, G.; Proc. Natl. Acad. Sci. U.S.A. 88, 854-8538, 1991
A;Title: The T-ceptor repertoire in the synovial fluid of a patient with rheumatch A;Reference number: A41299; MUID:92020887; PMID:1656449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; Species: Homo sapiens (man)
C; Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C; Accession: PH1771
R; Porcelli, S.; Yockey, C.E.; Brenner, M.B.; Balk, S.P.
J; Exp. Med. 178, 1-16, 1993
A; Title: Analysis of T cell antigen receptor (TCR) expression by human peripheral blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Pathway: heme biosynthesis; porphyrin biosynthesis
C;Superfamily: phytoene dehydrogenase
C;Keywords: heme biosynthesis; mitochondrion; oxidoreductase; porphyrin biosynthesis
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                                                    protoporphyrinogen oxidase (EC 1.3.3.4) - bovine (fragment)
C;Species: Bos primigenius taurus (cattle)
C;Date: 14-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 26-May-2000
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C;Date: 28-May-1992 #sequence_revision 28-May-1992 #text_change 05-Nov-1999
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A;Molecule type: mRNA
A;Molecule type: CUBA
A;Residues: 1-16 CUBA
A;Residues: 1-16 CUBA
A;Cross-references: GB:S57504; NID:g236332; PIDN:AAB19963.1; PID:g236333
C;Keywords: T-cell receptor
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50.0%; Pred. No. 4.7e+03;
tive 1; Mismatches 3; Indels
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Pred. No. 6.1e+03;
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Pred. No. 6.1e+03;
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28.6%;
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Best Local Similarity 36.*
Best Local 4; Conservative
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Matches 4; Conservative
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NTRTASKLTFG
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Best Local Similarity
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A;Molecule type: mRNA
A;Residues: 1-16 <POR>
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A; Molecule type: protein
A; Residues: 1-19 < ISH>
A; Residues: 1-19 < ISH>
A; Residues: 1-19 < ISH>
A; Consereferences: UNIPROT:Q9UCE9
A; Bxperimental source: Hela cell nuclei
A; Note: sequence extracted from NCBI backbone (NCBIP:134644)
C; Superfamily: ribonucleoprotein repeat homology
Query Match
Best Local Similarity 37.5%; Pred: No. 7.2e+03;
Matches 3; Conservative 4; Mismatches 1; Indels
Qy 4 RKSERIQR 11
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Db 8 KESERVDK 15
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Search completed: May 16, 2005, 13:07:13 Job time: 29.9231 secs

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0; Gaps

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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OM protein - protein search, using sw model

May 16, 2005, 12:39:11 ; Search time 85.7692 Seconds (without alignments) 89.556 Million cell updates/sec Run on:

US-08-869-386-1 77 1 RIQRGPGRAFVTIGK 15 Title: Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1612378 segs, 512079187 residues Searched:

16988 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

UniProt\_03:\*
1: uniprot\_sprot:\*
2: uniprot\_trembl:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

	Description	O10481 human immun	Q7zjt3 human immun	Q9e8s7 human immun	Q9enm9 human immun	Q8aqx9 human immun	Q8agy0 human immun	O9qex7 human 1mmun	Q6u2m7 sechium edu	Q8aqy1 human immun	Q8aqy2 human immun	homo	homo	homo	melea	gall		Q7mlw9 arabidopsis			Q6rck2 pseudomonas	Q6u2m9 momordica c		-		Q84038 influenza a					Q9pwq4 gallus gall	Q6v0x7 serratia ma
	ID	010481	Q7ZJT3	Q9E8S7	Q9ENM9	Q8AQX9	QBAQYO	Q9QEX7	Q6U2M7	QBAQY1	QBAQY2	Q9UCK9	OBUCLO	Q16228	Q6EML0	Q6EML1	Q924C7	Q7M1W9	Q6JCN3	Q95KS4	Q6RCK2	Q6U2M9	Q6TQT6	Q6U2N2	Q7M1V8	Q84038	Q9P2A2	080630	090633	Q7R974	Q9PWQ4	Q6V0X7
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de	Query	54.5	54.5	45.5	45.5	45.5	45.5	40.3		36.4	36.4	35.1	35.1	35.1	35.1	35.1	35.1	34.4	33.8	33.8	33.8	ω.	ë.	۳.	ä	32.5	32.5	ä	32.5	ä	32.5	32.5
	Score	42	42	35	35	35	35	31	29	28	28	27	27	27	27	27	27	26.5	26	26	56	56	26	56	25	25	25	25	25	25	25	25
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Q7MZM8 069173 069173 Q8UUH2 Q8UUH2 Q8UH4 Q9AH71 BLP4 BOMOR Q8JH97 Q9TRK1 COXO RAT Q7SOCS Q7PE81
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## ALIGNMENTS

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MEDLINE=20411423; PubMed=10954550;
MEJOLINE=20411423; PubMed=10954550;
DOI=10.1128/JVI.74.18.8494-8501.2000;
Melson J.A.E., Baribaud F., Edwards T., Swanstrom R.;
Patterns of changes in human immunodeficiency virus type 1 V3 sequence populations late in infection.";
J. Virol. 74:8494-8501(2000).
GO, GO:0019031; C:viral envelope; IEA.
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                                                                                                                                                                                                                                                                                                                                6108EAEC9C0CA947 CRC64;
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01-MAR-2003 (TrEWBLrel. 23, Created)
01-MAR-2003 (TrEWBLrel. 23, Last sequence update)
01-OCT-2003 (TrEWBLrel. 25, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Name=env;
Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBL_TaxID=11676;
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
                                                  Human immunodeficiency virus 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus
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              Envelope glycoprotein (Fragment).
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SEQUENCE
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                  MEDLINE=22439926; PubMed=12552446;

A REDLINE=2439926; PubMed=12552446;

I versen A.K.N., Christiansen C.B., Attermann J., Eugen-Olsen J.,

Schulman S., Berntorp E., Ingerslev J., Pugger L., Scheibel E.,

Tengborn L., Gerstoft J., Dickmeiss E., Svejgaard A., Skinhoj P.,

"Limited protective effect of the CCR5Delta32/CCR5Delta32 genotype on human immunodeficiency virus infection incidence in a cohort of patients with hemophilia and selection for genotypic X4 virus.";

J. Infect. Dis. 187:215-225 (2003).

EMBL; AX150666; AA061698.1;

GO; GO:0019028; C:nitegral to membrane; IEA.

GO; GO:0019031; C:viral capsid; IEA.

GO; GO:0019031; C:viral capsid; IEA.

GO; GO:0019031; C:viral molecule activity; IEA.

RO; GO:0019031; C:viral molecule activity; IEA.

RO; GO:005199; Fistructural molecule activity; IEA.

RO; GO:005199; Fistructural molecule activity; IEA.

REPER: PF000777; GP120.

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MEDLINE=20411423; PubMed=10954550;
DOI=10.1128/JVI.74.18.8494-8501.2000;
Melson J.A.E., Baribaud F., Edwards T., Swanstrom R.;
"Patterns of changes in human immunodeficiency virus type 1 V3 sequence populations late in infection.";
J. Virol. 74.8494-8501(2000).
BMBL; API55888; AAG09330.1;
GO; GO:0019031; C:viral envelope; IEA.
InterPro; IPR011056; Pept_S24_S26_C.
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Last annotation update)
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
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Last sequence update)
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Pred. No. 71;
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Pred. No. 4.2;
2; Mismatches
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Q9ENM9;
01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
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Matches 7; Conservative
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MEDLINE=22860939; PubMed=14502005; Freel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J., Freel S.A., Fiscus S.A., Pilcher C.B., Eron J.J. Jr., Shugars D.C.; Patrick B., Lennox J.L., Hicks C.B., Eron J.J. Jr., Shugars D.C.; Envelope diversity, coreceptor usage and syncytium-inducing phenotype of HIV-1 variants in saliva and blood during primary infection.", AIDS 17:2025-2033 (2003).
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GO, GO:0016021; C:integral to membrane; IEA.
GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0019031; C:viral envelope; IEA.
GO; GO:001908; F:seructural molecule activity; IEA.
InterPro; IPR000777; GP120.
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Envelope glycoprotein (Fragment).
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tes 6; Conservative
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MEDLINE=21103026; PubMed=11170057;

DOI=10.1002/1096-9071(200103)63:3<197::AID-JMV1000>3.3.CO;2-G;

Lin H.J., Siwak E.B., Lauder I.J., Hollinger F.B.;

Lin G. G. Glossylation sites.";

J. Med. Virol. 63:197-202(2001).

R. Med. Virol. 63:197-202(2001).

R. Med. Virol. 63:197-202(2001).

R. M. G. G. Glossylation sites.";

R. G. Glossylation sites.";

R. G. Glossylation sites.";

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MEDLINE-22860939; PubMed=14502005;

Preel S.A., Flacus S.A., Plicher C.D., Menezes P., Giner J.,

Patrick E., Lennox J.L., Hicks C.B., Bron J.J. Jr., Shugars D.C.;

"Envelope diversity, coreceptor usage and syncytium-inducing phenotype of HIV-1 variants in saliva and blood during primary infection.";

AIDS 17:2025-2031(203).

EMBL; AF536913; AAM63928.1; -.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0019029; C:viral capsid; IEA.

GO; GO:0019031; C:viral envelope; IEA.

GO; GO:0019031; C:viral envelope; IEA.

GO; GO:0019031; G:viral envelope; IEA.
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25 AA; 2749 MW; 9B6E9DACB8D56C0C CRC64;
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               01-WAR-2003 (TrEMBLrel. 23, Created)
01-WAR-2003 (TrEMBLrel. 23, Last sequence update)
01-0CT-2003 (TrEMBLrel. 25, Last annotation update)
Envelope glycoprotein (Fragment).
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
                                                                                                                         Viruses, Retroid viruses, Retroviridae, Lentivirus
NCBI_TaxID=11676;
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77.8%; Pred. No. 77;
tive 0; Mismatches
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program in the phloem of minor veins.", Plant Physiol. 133:1229-1239 (2003).

EMBL; AY379782; AAQ74884.1; -. GO, GO:0047216; F:inositcol 3-alpha-galactosyltransferase acti. . .; IEA. GO; GO:0016757; F:transferase activity, transferring glycosyl. . .; IEA. Glycosyltransferase; Transferase.

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MEDLINE=22860939; PubMed=14502005;

MEDLINE=22860939; PubMed=14502005;

Preel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,

Patrick E.A., Fiscus S.A., Pilcher C.B., Bron J.J. Jr., Shugars D.C.;

"Envelope diversity, coreceptor usage and syncytium-inducing phenotype of HIV-1 variants in saliva and blood during primary infection.";

ALIOS 17.2025-2033(2003).

EMBL; AR536912; AAM63927.1; -.

GO, GO:0019031; C:viral envelope; IEA.

InterPro, IPR000777; GP120.
                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Cucurbitales; Cucurbitaceae; Sechium.
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MEDLINE=22975109; PubMed=14526110; DOI=10.1104/pp.103.027714;
Ayre B.G., Blair J.B., Turgeon R.;
"Functional and phylogenetic analyses of a conserved regulatory
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Pred. No. 8.2e+02;
0; Mismatches 1; Indels
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05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Galactinol synthase (EC 2.4.1.123) (Fragment).
Name=GAS1;
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
VCBI_TaxID=11676;
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Pred. No. 1.4e+03;
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MEDLINE=94290510; PubMed=8019570;
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Hum. Mutat. 3:321-323(1994).
EMBL; 873627; AAB31191.1; -.
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Best Local Similarity 60.0%;
Matches 6; Conservative
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01-NOV-1996 (TrEMBLrel. 01,
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 5; Conservative
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                         4 RGPGRAF 10
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MEDLINE=22860939; PubMed=14502005;
MEDLINE=22860939; PubMed=14502005;
Freel S.A., Fiscus S.A., Pilcher C.D., Menezes P., Giner J.,
Patrick E., Lennox J.L., Hicks C.B., Eron J.J. Jr., Shugars D.C.;
Patrick E., Lennox J.L., Hicks C.B., Eron J.J. Jr., Shugars D.C.;
"Envelope diversity, coreceptor usage and syncytium-inducing phenotype of HIV-1 variants in saliva and blood during primary infection.";
AIDS 17:2025-2033(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=93099171; PubMed=1463770; DOI=10.1016/0925-4439(92)90068-X; Baba S., Takahashi T., Kasama T., Shirasawa H.; Baba S., Takahashi T., Kasama T., Shirasawa H.; Satahashi T., Kasama T., Shirasawa H.; Indentification of two novel amyloid A protein subsets coexisting in an individual patient of AA-amyloidosis."; Blochim. Blophys. Acta 1180:195-200(1992).

PIR, AZ790Z, YLHUA.

GO; GO:0005576; C:extracellular; IEA.

GO; GO:000553; P:actracelhalar; IEA.

InterPro; IRP000095; Serum_amyloid_A.
     Gaps
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 16, Last annotation update)
Serum amyloid A isotype 2 alpha protein (Serum amyloid A protein)
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Pred. No. 1.4e+03;
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Pred. No. 1.4e+03;
0; Mismatches 3; Indels
   Indels
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SEQUENCE 16 AA; 1612 MW; 1CAB4F077C9C8CC1 CRC64;
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                                                                                                                                           (TrEMBLrel. 23, Created)
(TrEMBLrel. 23, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
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71.4%;
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       6; Conservative
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                              2 IQRGPGRAF 10
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Best Local Similarity
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01-MAR-2003 (
01-OCT-2003 (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDINES-91099171; PubMed=1463770; DOI=10.1016/0925-4439(92)90068-X; Baba S., Takahashi T., Kasama T., Shirasawa H.; Baba S., Takahashi T., Kasama T., Shirasawa H.; Intakanashi T., Manasama T., Manasawa T., Manasama T., Ma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gape
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                                                                                            01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Serum amyloid A isotype 1 protein (Serum amyloid A protein)
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Last annotation update)
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16 AA.
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STRAIN-129/504;
MEDLINE-2129298B; PubMed=11262390; DOI=10.1074/jbc.M009382200;
Lovahin J.A., Estall J., Yusta B., Brown T.J., Drucker D.J.;
Lovahin J.A., Estall J., Yusta B., Brown T.J., Drucker D.J.;
"Glucagon-like peptide (GLP)-2 action in the murine central nervous
system is enhanced by elimination of GLP-1 receptor signaling.";
Biol. Chem. 276:21489-21499 (2001).
EMBL; AF338224; AAK63043.1;
MGD; MGI:2136733; Glp2r.
GO; GO:0016021; C:integral to membrane; TAS.
GO; GO:0016021; C:integral to membrane; TAS.
GO; GO:0016021; P:glucagon receptor activity; TAS.
GO; GO:000186; P:G-protein coupled receptor protein signalin. ..; TAS.
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01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Porin porl (Fragment).
Arabidopsis thaliana (Mouse-ear cress).
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyjedons; core eudicots; rosids;
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
NCBI_TAXID=3702;
Gaps
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                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID=10090;
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Pred. No. 1.5e+03;
2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 27; DB 2; Length 22;
Pred. No. 1.9e+03;
3; Indels
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22 AA; 2526 MW; 2C5BF53DCCD425C9 CRC64;
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                                                                                                                                                                                                                                                                   01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Glucagon-like peptide-2 receptor (Fragment).
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Submitted (JUL-1994) to the PIR data bank.
PIR; PA0045; PA0045.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14 AA.
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  Mismatches
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7
                                                2 IQRG---PGRAFV-TIG 14
                                                                                              1 iQTGVDNPGHPFIMTVG 17
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14 AA; 1546 MW;
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Best Local Similarity 50.v.
Best Local 6; Conservative
     8; Conservative
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Best Local Similarity
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01-DEC-2001 (
01-DEC-2001 (
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Q924C7
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SEQUENCE FROM N.A.

Axelsepon E., Smith N.G., Sundstrom H., Berlin S., Ellegren H.,

Axelsepon E., Smith N.G., Sundstrom H., Berlin S., Ellegren H.,

"Male-blassed mutation rate and divergence in autosomal, z-linked and

"Alinked introns of chicken and Turkey.",

Mol. Biol. Evol. 21:1538-1547(2004).

EMBL, AX13962, AAN74866.1, -

EMBL, AX13962, AAN74866.1, -

EMBL, AX13962, Fitnansferase activity, IEA.

GO, GO:0016772; Fitransferase activity, transferring phosphor. . .; IEA.

InterPro; IPR000749; ATP-gua Ptrans.

Pfam; PF02807; ATP-gua Ptrans.
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Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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                                                                                                                                                                                                                                                                Meleagris gallopavo (Common turkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Meleagris.
                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=15140948; DOI=10.1093/molbev/msh157;
Axelsson E., Smith N.G., Sundstrom H., Berlin S., Ellegren H.;
Male-biased mutation rate and divergence in autosomal, z-linked in w-linked introns of chicken and Turkey.";
Mol. Biol. Evol. 21:1538-1547(2004).
BMBL; AX139863; AAN74867.1; -..
GO; GO:0016702; Fitname activity; IEA.
GO; GO:0016702; Fitname activity; transferring phosphor...
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                                                                                                                                                                                                                                              B-creatine kinase (Fragment).
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47.1%;
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NCBI_TaxID=9940;
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                                                                                                                                                                  SEQUENCE
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Q6U2M9
                                                                                                                                                                                                                                              Matches
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SEQUENCE FROM N.A.
STRAIN=DAEC18, DAEC19, DAEC214, DAEC7, DAEC5, DAEC20,
STRAIN=DAEC18, DAEC120, DAEC114, DAEC11b, DAEC18, DAEC19, EC7372,
BCOR37, DAEC9, ECOR50, IH1128, C1845, and ECOR64;
PubMed=15014151; DOI=10.1093/molbev/msh118;
BSCOber-Paramo P. (Iermont O., Blanc-Potard A.B., Bui H.,
Le Bouguenec C., Denamur E.;
"A Specific Genetic Background Is Required for Acquisition and
Expression of Virulence Factors in Escherichia coli.";
Mol. Biol. Evol. 21:1085-1084(2004).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name-dāp12;
Ovis aries (Sheep).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
Caprinae; Ovis.
                                                                                                                                                                                                                                              Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 26; DB 2; Length 20;
Pred. No. 2.6e+03;
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                                                                                                                               06JCN3;
05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Last sequence update)
Last annotation update)
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                                                                                                                 20 AA
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                                                                                                                 PRT;
                                                                                                                                                                                                                                                            Enterobacteriaceae; Escherichia
NCBI_TaxID=562;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; AYS25516; AAT00552.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL, AY525528; AAT00576.1; -
EMBL, AY525529; AAT00578.1; -
EMBL, AY525530; AAT00580.1; -
EMBL, AY525531; AAT00582.1; -
EMBL, AY525532; AAT00584.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL, AY525517; AAT00554.1;
EMBL, AY525518; AAT00556.1;
EMBL, AY525519; AAT00568.1;
EMBL, AY525520; AAT00560.1;
EMBL, AY525521; AAT00564.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33.8%;
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 2 KGPG-LYTEIGK 12
                                                                                                               PRELIMINARY;
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SEQUENCE
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EMBL;
EMBL;
EMBL;
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EMBL;
EMBL;
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                                                                            RESULT 18
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EMBL; AY504985; AAS46984.1; -.
GO; GO:0016702; F:oxidoreductase activity, acting on single d. . .; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Momordica charantia (Bitter gourd) (Balsam pear).
Momordica, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Wagnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Cucurbitales; Cucurbitaceae; Momordica.
NCBI_TaxID=3673;
                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=MT15;
Hendrickx B., Junca H., Vosahlova J., Faber F., Lindner A., Ruegg Bucheli-witschel M., Egli T., Mau M., Schlomann M., Brennerova M., Brenner V., Pieper D., Top E., Dejonghe W., Bastiaens L., Springael D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria, Proteobacteria, Gammaproteobacteria, Pseudomonadales,
Pseudomonadaceae, Pseudomonas.
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MEDLINE=22975109; PubMed=14526110; DOI=10.1104/pp.103.027714;
                                                                                                                                                                                                               Score 26; DB 2; Length 21;
Pred. No. 2.7e+03;
1; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33.8%; Score 26; DB 2; Length 21; 50.0%; Pred. No. 2.7e+03;
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SEQUENCE FROM N.A.
TISSUB-Peripheral blood;
TISSUB-Peripheral blood;
Subils S.A., Staines K.A.;
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ419229; CAD11671.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21 21
21 AA; 2191 MW; 4200125E64F0F0DE CRC64;
                                                                                                                                                                    BE2E264A4CD38D6D CRC64;
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Galactinol synthase (EC 2.4.1.123) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-JUL_2004 (TrEMBLrel. 27, Created)
05-JUL_2004 (TrEMBLrel. 27, Last sequence update)
05-JUL_2004 (TrEMBLrel. 27, Last annotation update)
Catechol 2,3-dioxygenase (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21 AA.
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                                                                                                                                                                                                                    33.8%;
54.5%;
                                                                                                                                                                       21 AA; 2316 MW;
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Matches 7; Conservative
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                                                                                                                                                                                                                                        Local Similarity 54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                2 IQRGPGRAFVT 12
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S VPRGRGAAEVT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pseudomonas putida.
                                                                                                                               21
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NON TER
NON TER
SEQUENCE 21
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GO:0047216; F:inositol 3-alpha-galactosyltransferase acti. GO:0016757; F:transferase activity, transferring glycosyl.
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Nucleic Acids Res. 6:3745-3757(1979)
EMBL; J02114; AAA43398.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAR-2004 (TrEMBLrel. 26, Created)
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    GO, GO:0047216; F:inositol 3-alphe
GO; GO:0016757; F:transferase acti
Glycosyltransferase; Transferase.
NON TER
SEQÜENCE 24 AA; 2538 MW; E2BCC
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Local Similarity 71.4%;
hes 5; Conservative
                                                                                                                                                                                                                                                          33.8%;
                                                                                                                                                                                                                                                                                            Best Local Similarity 71.4%;
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                              6 PGRAFVT 12
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GRSFVPI 9
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NCBI_TaxID=11320;
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SEQUENCE
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Q7M1V8;
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Matches
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Q84038
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
eurosids I; Cucurbitales; Cucurbitaceae; Citrullus.
Ayre B.G., Blair J.E., Turgeon R.;
"Functional and phylogenetic analyses of a conserved regulatory program in the philoem of minor veins.";
Plant Physiol. 133:1229-1239 (2003).

EMBL; AY379780; AAQ74882.1; -..

GO; GO:0047216; F:inositol 3-alpha-galactosyltransferase acti...

GO; GO:016757; F:transferase activity, transferring glycosyl...

Glycosyltransferase; Transferase.

NON TER 23

SEQÜENCE 23 AA; 2444 MW; 62411699CAB81657 CRC64;
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 23;
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Kennedy M.C., Dietrich F.S.;
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AY38930.2 AAQ97234.1; -.
NON TER 1 1 SEQÜENCE 24 AA; 2866 MW; 83820AB41EF59E7C CRC64;
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Galactinol synthase (EC 2.4.1.123) (Fragment).
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 26; DB 2;
Pred. No. 3e+03;
1; Mismatches
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Plant Physiol. 133:1229-1239(2003).
EMBL; AX379777; AAQ74879.1; -.
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Matches 5, Conservative
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16 PKRAYVT 22
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Name=YHR065C;
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Matches 5; Conserv
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledona; core eudicots; asterids;
lamiids; Solanales; Solanaceae; Nicotiana.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bauw G., De Loose M., Inze D., Van Montagu M., Vandekerckhove J.;
"Alterations in the phenotype of plant cells studied by NH2-terminal
amino acid-sequence analysis of proteins electroblotted from two-
dimensional gel-separated total extracts.";
Proc. Natl. Acad. Sci. U.S.A. 84:4806-4810(1987).
PIR. D28027, D28027.
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-UDN-2003 (TrEMBLrel. 24, Last annotation update)
Influenza Afpv/rostock/34 (h7nl), neuraminidase (seg 6), 3' end vrna (initiator region for protein coding). (Fragment).
Influenza A virus.
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Pred. No. 2e+03;
1; Mismatches 1; Indel8
                                                            Length 24;
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                                                                                                                      1; Indels
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E2BC0AE9D7930C06 CRC64;
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60; GO:0016020; C:membrane; IEA.
GO; GO:0016020; F:exc-alpha-sialidase activity; IEA.
GO; GO:0004308; F:exc-alpha-sialidase
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01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Protein P7 (Fragment).
                                                         Score 26; DB 2; 1
Pred. No. 3.1e+03;
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32.5%;
62.5%;
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Best Local Similarity 62.5%;
Matches 5; Conservative
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Best Local Similarity 62.5
Matches 5; Conservative
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Q7R974
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Nishizawa M., Nakajima T., Yasuda K., Kanzaki H., Sasaguri Y.,
Watanabe K., Ito S.,
"Close kinship of human 20alpha-hydroxysteroid dehydrogenase gene with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
Alphaherpesvirinae; Simplexvirus.
NCBI_TaxID=36347;
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MEDLINE=98440589; PubMed=9765470;
Smith A.L., Black D.H., Eberle R.;
"Molecular evidence for distinct genotypes of monkey B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 14;
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NiBhizawa M., Nakajima T., ItO S.;
Submitted (FBE-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AB037903; BAA9288811; -.
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   InterPro; IPR001860; Glyco_hydro_34.
Pfam; PF00064; Neur; 1.
NON TER 12 12 12 SEQÜENCE 12 AA; 1316 MW; DC0B3CE899505326 CRC64;
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Pred. No. 2.8e+03;
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                                                                                                                                                                                                                                                                    2; Mismatches
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01-0CT-2000 (TrEMBLrel. 15, Last sequence
01-0CT-2000 (TrEMBLrel. 15, Last annotation of the control of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 three aldo-kero reductase genes.";
Genes Cells 5:111-125(2000).
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80.0%;
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                                                                                                                                                                                                 32.5%;
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Les 4; Conservative
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                                                                                                                                                                                                                            Best Local Similarity 44.4
Matches 4; Conservative
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9 PGRSF 13
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090630
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Q7M2M8;
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                                               "Genome sequence and comparative analysis of the model rodent malaria
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                    Gaps
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Salzberg S.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M.J.,
Carucci D.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Plasmid pSMT1.
Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
                                                                parasite Plasmodium yoelii yoelii.",
Nature 419:512-519(2002).
-!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
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MEDLINE=20078374; Pubmed=10612250;

MEDLINE=20078374; Pubmed=10612250;

Mido Y., Burt D.W., Paton I.R., Sharp P.J., Dunn I.C.;

"Mapping of the prolactin gene to chicken chromosome 2.";

Anim. Genet. 30:473-473(1999).

EMBL; AJ239131; CAB43530.1; -.

EMBL; AJ239131; CAB43530.1; -.

EMBL; AJ239131; CAB43530.1; -.

NON TER 20 20

SEQÜENCE 20 AA; 2223 MW; 258CC8CAA95F12D6 CRC64;
                                                                                                                                                                                                                                                                       32.5%; Score 25; DB 2; Length 20; 40.0%; Pred. No. 3.9e+03; tive 3; Mismatches 3; Indele
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Pred. No. 3.9e+03;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                 20 AA; 2461 MW; C583B1AD3B45C3FC CRC64;
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01-WAY-2000 (TrEMBLrel. 13, Last sequence update)
01-WAR-2003 (TrEMBLrel. 23, Last annotation update)
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(TrEMBLrel. 27, Last sequence update)
(TrEMBLrel. 27, Last annotation update)
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                                                                                                                                                                                                                                                                                                                3; Mismatches
                                                                                                                                             preliminary data.
EMBL; AABLO1002466; EAA19325.1; -.
Hypothetical protein.
NON TER
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NCBI_TaxID=615;
                                                                                                                                                                                                                                                                                             Local Similarity 40.0 tes 4; Conservative
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Matches 4; Conservative
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7 GRGFIT 12
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05-JUL-2004
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09 PW04;
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Matches
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Q9PWQ4
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Q6V0X7
      STERRES
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MEDLINE=90354445; PubMed=2167319;
MEANMATUILLAM M., Radke G.A., Andrews P.C., Roche T.E.;
Meanges in the core of the mammalian-pyruvate dehydrogenase complex upon selective removal of the lippyl domain from the transacetylase component but not from the protein X component.";
J. Biol. Chem. 265:14512-14517(1990).
PIR, A37823; A37823.
PIR, A37823; A37823.

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NON TER 17 17
NON TER 17 17
SEQÜENCE 17 AA; 1743 MW; 5BFC5FB662D014D5 CRC64;
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Cetartiodactyla, Ruminantia, Pecora, Bovidae,
Bovinae, Bos.
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Bacteria, Proteobacteria, Gammaproteobacteria, Enterobacteriales,
Enterobacteriaceae, Yersinia.
NOBI_TaxID=632;
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01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last amnotation update)
Dihydroliposmide S-acetyltransferase (EC 2.3.1.12) (Pragment).
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Pred. No. 4.1e+03;
1; Mismatches 1; Indels
                                                                                                                                                                                                 Length 22;
                                                                                                                                                                                                                                                     1; Indels
Manzoor S.E., Gill M.J., Thomas C.M.;
Submitted (JUL-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AY355287; AAR18691.1; -.
Plasmid.
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Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases
EMBL; AF065382; AAC17437.1; -.
NON_TER 15 15
                                                                                                                    22 22
22 AA; 2039 MW; 8F6B89343C822704 CRC64;
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    Last sequence update)
    Last annotation update)

                                                                                                                                                                                           'Match 32.5%; Score 25; DB 2; I Local Similarity 66.7%; Pred. No. 4.3e+03; les 4; Conservative 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17 AA
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Putative ferrochelatase HemH (Fragment).
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01-AUG-1998 (TrEMBLrel. 07, Last seq
01-DEC-2001 (TrEMBLrel. 19, Last ann
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66.7%;
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hes 6, Conservative
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4 RGPGRAF 10
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Best Local Similarity
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                            RESULT 36
Q9R4H4
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Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 24; DB 2; Length 19;
Pred. No. 5.6e+03;
1; Mismatches 2; Indels
   Length 15;
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Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF459839; AAL67519.1; -.
                                                           6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Li H., Deeb N., Zhou H., Ashwell C.M., Lamont S.J.,
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF459837; AAL67517.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NON TER 1 1
NON TER 19 19
SEQUENCE 19 AA, 2046 MW, 1250C1CBFE03C2F7 CRC64;
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19 AA; 2032 MW; 1315C1CBFE03C2F7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAR-2002 (TrEMBLrel. 20, Last sequence update) 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Score 24; DB 2; 1
Pred. No. 4.5e+03;
2; Mismatches 6;
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         31.2%;
38.5%;
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57.1%;
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2 MQSKPGVLMVNLG 14
                                                                                                                              2 IQRGPGRAFVTIG 14
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                                                                  5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TGF-beta4 (Fragment).
Gallus gallus (Chicken).
                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 EMGPGHA 10
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SEQUENCE
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Matches
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Q8UVE
1D Q8UVE
AC Q8UVE
DT 01-MA
DT 01
                                                                        Matches
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MEDLINE=21116988; PubMed=11179344;
MEDLINE=21116988; PubMed=11179344;
MOI-10.1128/IAI.69.3.1687-1696.2001;
MAIDER C.M., Blum E., Miller Y.K., Ryan D., Popovic T., Stephens D.S.;
"exl, an exchangeable genetic island in Neisseria meningitidis.";
"Infect. Immun. 69:1687-1696(2001).
NON_TER I.
NON_TER I.
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                                                                                                                                                                                                                                                                                                   MEDLINE-96085152; PubMed-8521853;
Steuber J., Arendsen A.F., Hagen W.R., Kroneck P.M.;
Steuber J., Arendsen A.F., Hagen W.R., Kroneck P.M.;
Molecular properties of the dissimilatory sulfite reductase from "Molecular properties of the dissimilatory sulfite reductase from Desulfovibrio desulfuricans (Essex) and comparison with the enzyme from Desulfovibrio vulgaris (Hildenborough).";
Fur. J. Biochem. 23:873-879(1995).
PIR; 863490, 563490.
GO; GO:0018551; F. hydrogensulfite reductase activity; IEA.
SEQUENCE 20 AA; 2193 MW; F939E03B6E355135 CRC64;
                                                                                                                                                                       Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales; Desulfovibrionaceae; Desulfovibrio.
                                   01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-0CT-2003 (TrEMBLrel. 25, Last annotation update)
Sulfite reductase 50 kDa alpha subunit (EC 1.8.99.3) (Fragment).
Desulfovibrio desulfuricans.
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Bacteria; Proteobacteria; Betaproteobacteria; Neisseriales;
Neisseriaceae; Neisseria.
Nois TaxID=487;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 24; DB 2; Length 22;
Pred. No. 6.5e+03;
1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 24; DB 2; Length 20;
Pred. No. 5.9e+03;
4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE 22 AA; 2584 MW; FIBEBC6F2F3C2C49 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Created)
Last sequence update)
Last annotation update)
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20 AA.
PRT;
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BLP4_BOWOR
ID BLP4_BOWOR
STANDARD;
AC P29005;
DT 01-DEC-1992 (Rel. 24, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        h 31.2%;
Similarity 57.1%;
4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 36.4
Matches 4; Conservative
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  PRELIMINARY;
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GO; GO:0016301; F:kinase activity; IEA.
GO; GO:0016772: F:transferase activity, transferring phosphor. . .; IEA.
InterPro; IPR000749; ATP-gua_Ptrans.
Pfam; PF02807; ATP-gua_PtransN; 1.
                                                                                                                                                                                                                                                                                  Bures S., Nadvornik P., Saetre G.-P.; "Hybridization and apparent hybridization between meadow pipit (Anthus pratensis) and water pipit (A. spinoletta).";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Production of anti-NC1 antibody by affected male dogs with X-linked hereditary nephritis: a probe for assessing the NC1 domain of collagen type IV in dogs and humans with hereditary nephritis."; Submitted (FEB-1993) to the BMEL/GenBank/DDBJ databases. SEQUENCE 21 AA; 2300 MW; 08C6D4D9DJD62EEA CRC64;
                                                                                                                                                                          Eukaryoča; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Passeriformes; Motacillidae; Anthus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Merazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis.
NCBI_TaxID=9615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thorner P.S., Baumal R., Valli V.E., Mahuran D., Marrano P.M., Jacobs R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30.5%; Score 23.5; DB 2; Length 21; 66.7%; Pred. No. 7.6e+03; tive 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 23.5; DB 2; Length 16;
Pred. No. 5.9e+03;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16 AA; 1726 MW; 106D0486800C21E7 CRC64;
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25-CGT-2004 (TrEMBLrel. 28, Last annotation update)
Collagen type IV 24 kDa component (Fragment)
Canis familiaris (Dog).
                                                               01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Created)
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EMBL; AF527052; AAM93207.1; -.
HSSP; P05122; 1QH4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30.5%;
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nes 6, Conservative
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                           PRELIMINARY;
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                             SEQUENCE FROM N.A
                                                                                                                                                                                                                  NCBI_TaxID=45803;
                                                                                                                                                       Anthus pratensis.
                                                                                                                                     B-CK (Fragment
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ID COXO RAT
AC P80432;
DT 01-NOV-199
DT 01-NOV-199
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                           Q8JH97
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Q9TRK1
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  Q8JH97
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                                                                                                                                                                                         TISSUE=Skin secretion;

X MEDLINE=92078177; PubMed=1744108;
A Gibson B.W., Tang D., Mandrell R., Kelly M., Spindel B.R.;
A Gibson B.W., Tang D., Mandrell R., Kelly M., Spindel B.R.;
A Gibson B.W., Tang D., Mandrell R., Kelly M., Spindel B.R.;
T "Bombinin-like peptide motion orientalis.";
I. J. Biol. Chem. 266:21103-23111(1991).
- I- FUNCTION: Has antimidicrobial activity, but no hemolytic activity.
C -1- FUNCTION: Has antimidicrobial activity, but no hemolytic activity.
C -1- SUBCELLULAR LOCATION: Secreted.
- I- SIBSUE SPECIFICITY: Skin.
- I- SIMILARITY: Belongs to the bombinin family.
C -1- SIMILARITY: Belongs to the bombinin family.

R PIR; D41575; D41575.
W Amidation; Amphibian defense peptide; Antibiotic;
W Direct protein sequencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GO; GO:0016301; F:kinase activity; IEA.
GO; GO:0016772; F:transferase activity, transferring phosphor. . .; IEA.
InterPro; IPR000749; ATP-qua_Ptrans
Pfam; PF02807; ATP-qua_PtransN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bures S., Nadvornik P., Saetre G.-P.;
"Hybridization and apparent hybridization between meadow pipit (Anthus pratensis) and water pipit (A. spinoletta).";
Herediza 13:0-0(2002).
EMBL; AFS27053; AAM93208.1; -.
01-DEC-1992 (Rel. 24, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Bombinin-like peptide 4 (BLP-4).
Bombina orientalis (Oriental fire-bellied toad).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Amphibia; Batrachia; Anura; Archeobatrachia; Bombina.
NCBI_TaxID=8346;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Archosauria, Aves, Neognathae, Passeriformes, Motacillidae, Anthus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 24; DB 1; Length 25;
Pred. No. 7.3e+03;
2; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25 25 Phenylalanine amide.
25 AA; 2409 MW; E97916634BC3F768 CRC64;
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anthus spinoletta (Water pipit).
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Matches 5; Conserv
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98JH96

RESULT 39

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Best Loc Matches

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EMBL, AABX01000510; EAA28761.1; -.
SEQUENCE 11 AA; 1251 MW; 48F2534E31B2C9C3 CRC64;
                                        Query Match
Best Local Similarity 80.07
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Best Local Similarity
5, Conserva
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ENSANGP00000024647
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                                                                                                                                             10 FVTIG 14
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Zea mays (Maize)
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Q7PE81;
01-MAR-2004 (
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P80625;
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UC19 MAIZE
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    -!- CATALYTIC ACTIVITY: 4 ferrocytochrome c + 0(2) = 4 ferricytochrome

                                                                                          Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
      05-JUL-2004 (Rel. 44, Last annotation update)
Cytochrome c oxidase polypeptide VIIc, mitochondrial (BC 1.9.3.1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is preliminary data.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       -i- SIMILARITY: Belongs to the cytochrome c oxidase VIIc family. PIR; S65388; S65388.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Neurospora crassa.
Bukaryota, Fungi, Ascomycota, Pezizomycotina, Sordariomycetes,
Sordariomycetidae, Sordariales, Sordariaceae, Neurospora.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 23; DB 1; Length 10;
Pred. No. 4.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein sequencing; Inner membrane; Mitochondrion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NON TER 10 10 10 SQUENCE 10 AA; 1117 MW; 126DE767687B1B0B CRC64;
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Last annotation update)
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                                                   (VIIIA) (Fragment).
Name=Cox7c; Synonyms=Cox7c1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAR-2004 (TrEMBLrel. 26, 01-MAR-2004 (TrEMBLrel. 26, 01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Conservative
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4 EEGPGK 9
                                                                                                                                             NCBI_TaxID=10116;
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C750C5
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Touzet P., Ricoardi F., Morin C., Damerval C., Huet J.-C.,
Pernollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein database: towards an integrated
                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Anopheles gambiae str. PEST.
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Anopheles.
NCBI_TaxID=180454;
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Theor. Appl. Genet. 93:997-1005 (1996).
-!- MISCELLANBOUS: On the 2D-gel the determined pI of this unknown protein is: 5.6, its MW is: 18.4 kDa.
Maize-2DPAGE; P80625; COLEOPTILE.
MaizeDB; 123951; -.
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anopheles Genome Sequencing Consortium;
Submitted (APR-2003) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
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01-OCT-1996 (Rel. 34, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Unknown protein from 2D-PAGE of etiolated coleoptile (Spot 406)
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Pred. No. 6.3e+03;
Length 11;
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                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        preliminary data.
EMBL; AAAB01004344; EAA45843.1; -.
SEQUENCE 14 AA; 1652 MW; 4A8A0A1AEC3F7FD3 CRC64;
                                                                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation update)
Score 23; DB 2;
Pred. No. 5e+03;
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                                                                                                                                                                                                                                                                            14 AA.
                                                1; Mismatches
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29.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                       Name=ENSANGG00000020916;
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                                                                  Query Match
29.9%; Score 23; DB 1; Length 15;
Best Local Similarity 50.0%; Pred. No. 6.8e+03;
Matches 4; Conservative 1; Mismatches 3; Indels
NON TER 15 15 SEQÜENCE 15 AA; 1672 MW; 1CF69D4DA8737F9D CRC64;
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7 GRAFVTIG 14 || : | | 2 GRRYTTYG 9 ò g Search completed: May 16, 2005, 13:00:26 Job time : 88.7692 secs

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Sequence 17, A
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Sequence 2, Al
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Sequence 13,
Sequence 28,
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(cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

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          GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-08-318-025A-17

US-08-109-047-7

US-08-109-047-7

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US-08-109-06-7

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Maximum DB
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No.
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US-08-481-9858-35
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US-08-39-48-082-28
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US-08-482-388-3 US-07-950-571A-1 US-08-266-448-1 US-08-485-324-31 US-08-447-506-13 US-08-447-506-31 US-08-235-437-31 US-08-235-437-31 US-08-247-515-13

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.257 -460	-463	-464	-467	-493	-279	-488	1894-	-105	-709	-458	-657	-657	-657	594-	1894-	-657	1594-	1-657	1594-	-111	1-211	1593-	3-704	1594-	JS94-	-920	3-466 3-688	1892-	3- / U4 JS94-	7-920 7-466	9-688	3-257 3-460	3-463	3-464	3-462	3-467	JS94-	3-257 8-460	3-463	8-465 8-464	8-462	8-467 8-218	US92-	9-820
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47-51 93-87 55-62 55-68	60-98	55-62	55-68	55-68	86-09	4-051	55-62	55-66	4-051	5-03	04-17	55-62	55-62	55-68	55-66	60-98	6-09	4-026	4-05	4-05	9-09	63-96	64-3	62-5	2-06	55-6	55-6	55-6	55-6	55-6	55-6	55-6	6-09	6-090	6-090	6-090	160-9 14-05	94-05	74-05 94-05	94-05	34 - UD 155 - 6	155-6	34-05	279-9
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Sequence 7, Application US/08709047

Sequence 7, Application US/08709047

Sequence 7, Application US/08709047

Sequence 7, Application US/08709047

Parent NO. 5652333

SEMERAL INFORMATION:

APPLICANT: Fung, Michael S.C., Sun, Bill N.C, Sun, Cecily R.Y., Kim, Young W., Yu, APPLICANT: Liming

TITLE OF INVENTION: THE GCIQ RECEPTOR, HIV-1 gpl20 REGION BINDING THERETO,

TITLE OF INVENTION: AND RELATED PEPTIDES AND TARGETING ANTIBODIES

NUMBER OF SEQUENCES: 13

CORRESPONDENCE ADDRESS:

ADDRESSEB: Tanox Blosystems, Inc.

STREET: 10301 Stella Link Rd.

CONTRY: Houston

STATE: Texas

COUNTRY: USA
Methods and Compositions for Diagnosing
and Treating Certain HIV Infected Patients
197
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                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppatible
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 07/891,451
FILING DATE: 29-MAY-1992
ATTORNEY/AGBNT INFORMATION:
NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: 31,215
REFERENCE/DOCKET NUMBER: MST33A
TELECOMMUNICATION INFORMATION:
TITLE OF INVENTION: Methods and Compositions for District OF INVENTION: and Treating Certain HIV Infects NUMBER OF SEQUENCES: 197
CCRRESPONDENCE ADDRESS:
ADDRESSEE: Howson and Howson STREET: P.O. Box 457, 321 No. 5556744ristown Road CITY: Spring House STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 77; DB 1; I
100.0%; Pred. No. 8.5e-06;
tive 0; Mismatches 0;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/709,047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (215) 540-9206
TELEPAX: (215) 540-5818
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PS/2
OPERATING SYSTEM: DOS 3.30
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Best Local Similarity 100.
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         Sequence 51, Appl
Sequence 68, Appl
Sequence 16, Appl
Sequence 1, Appli
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                                                                                                                                                                                                                                                                                                                         APPLICANT: Leclerc, Claude
APPLICANT: Leclerc, Claude
APPLICANT: Sebo, Peter
APPLICANT: Sebo, Peter
APPLICANT: Sebo, Peter
APPLICANT: OF INVENTION: Recombinant Mutants for Inducing
TITLE OF INVENTION: Specific Immune Responses
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner
STREET: 1300 I Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: D.C.
COWTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/336,087
FILING DATE:
    US-09-454-204A-51
US-09-454-204A-68
US-09-508-552-16
PCT-US92-01303-1
                                                                                                                                           ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/011,644
FILING DATE: 29-JAN-1993
ATTORNEY AGENT INFORMATION:
NAME: Meyers, Kenneth J.
REGISTATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 03495-0109-TELEPHONE: 202-408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17, Application US/08218025A
Patent No. 5556744
GENERAL INFORMATION:
APPLICANT: Weiner, David B.
APPLICANT: Ugen, Kenneth E.
APPLICANT: Williams, William V.
                                                                                                                                                                                                                                    US-08-336-087-2
; Sequence 2, Application US/08336087
; Setent No. 5503829
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 RIORGPGRAFVTIGK 15
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MOLECULE TYPE: peptide
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Best Local Similarity
Matches 15; Conserv
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US-08-218-025A-17
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                                                                                                                                                                                                                                                                                                                                            DB 1; Length 15;
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APPLICANT: Ladant, Daniel
APPLICANT: Leclerc, Claude
APPLICANT: Sebo, Peter
APPLICANT: Ullmann, Agnes
TITLE OF INVENTION: Recombinant Mutants for Inducing
TITLE OF INVENTION: Specific Immune Responses
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: D.C.
COUNTRY: USA
COUNTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: PELOPS MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/479,400
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Meyers, Kenneth J.
REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 03495-0109-01000
                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                              100.0%; Score 77;
100.0%; Pred. No. 8
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APPLICATION NUMBER: US 08/011,644
FILING DATE: 29-JAN-1993
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US/08/410,360
      ATCOLNG DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Mirabel, Eric P.
REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TRX9:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION (713) 664-228
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TELEPHONE: 202-408-4000
TELEFAX: 202-408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 1300 I Street, N.W. CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIORGPGRAFVTIGK 15
                                                                                                                                                    TELEPHONE: (713) 664-220
TELEFAX: (713) 664-8914
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                         15 amino acids
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.0
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , MOLECULE TYPE: peptide US-08-479-400-2
                                                                                                                                                                                                                                                       ; TYPE: amino acid
; TOPOLOGY: linear
US-08-709-047-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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100.0%; Score 77; DB 1; Length 15;

Query Match

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APPLICANT: Liming
TITLE OF INVENTION: THE GCIQ RECEPTOR, HIV-1 gpl20 REGION BINDING THERETO,
TITLE OF INVENTION: AND RELATED PEPTIDES AND TARGETING ANTIBODIES
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Tanox Blosystems, Inc.
STREET: 10301 Stella Link Rd.
CITY: Houston
STARET: USA
COUNTRY: USA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
Fatent No. 5711947
GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A.
APPLICANT: Berzofsky, Jay A.
TITLE OF INVENTION: METHOD TO INDUCE CYTOTOXIC T LYMPHOCYTES
TITLE OF INVENTION: SPECIFIC FOR A BROAD ARRAY OF HIV-1 ISOLATES USING HYBRID TITLE OF INVENTION: SYNTHETIC PEPTIDES
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
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                        Gaps
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100.0%; Pred. No. 8.5e-06;
tive 0; Mismatches 0;
Best Local Similarity 100.0%; Pred. No. 8.5e-06; Matches 15; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch
COMPUTER: IBM PS/2
OPPRATING SYSTEM: DOS 3.30
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/410,360
FILING DATE:
CLASSIFICATION 530
PRIOR APPLICATION 530
FILING DATE:
APPLICATION DATA:
APPLICATION TOWNER:
FILING DATE:
APPLICATION TOWNER:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
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REGISTRATION NUMBER: 31,211
REFRENCE/DOCKET NUMBER: 1NX95-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (713) 664-8914
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 15 aning acids
                                                                                                                                                                                                                                                         Sequence 7, Application US/08410360 Patent No. 5691447 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                           1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-410-360-7
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Gaps
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APPLICANT: Kim, Young W., Yu, Liming
TITLE OF INVENTION: THE ECLG RECEPTOR, HIV-1 gpl20 REGION BINDING
TITLE OF INVENTION: ATTIENDES
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Tanox Blosystems, Inc.
STREET: 10301 Stella Link Rd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
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100.0%; Pred. No. 8.5e-06;
                                                                                                                                                                                      SUFTWARE: WOORDERIECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/707,801A
FILING DATE: 09/04/1996
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/410,360
FILING DATE: 03/24/1995
ATTORNEY/AGENT INFORMATION:
NAME: Mirabel, Eric P.
REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TNX95-1AA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (713) 664-2288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Houston
STATE: Texas
CONTYRY: USA
ZIP: 77025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch
COMPUTER: IBM PS/2
OPERATING SYSTEM: DOS 3.30
SOCTWARE: Wordberfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/709,006
FILING DATE: 09-SEP-1996
CLASSIFICATION NUMBER: 08/410,360
FILING DATE: 24-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: MITABLE, ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 7, Application US/08709006
Patent No. 5731428
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REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TN
TELECOMMUNICATION INFORMATION:
TELEPHONE: (713) 664-2288
                                                                                                   COMPUTER: IBM PS/2
OPERATING SYSTEM: DOS 3.30
SOFTWARE: Wordperfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (713) 664-8914
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPX: (713) 664-8914
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-707-801A-7
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APPLICANT: Fung, Michael S.C., Sun, Bill N.C, Sun, Cecily R.Y., Kim, Young W., Yu, APPLICANT: Liming
APPLICANT: Liming
TITLE OF INVENTION: THE GCIG RECEPTOR, HIV-1 gpl20 REGION BINDING THERETO,
TITLE OF INVENTION: AND RELATED PEPTIDES AND TARGETING ANTIBODIES
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Tanox Blosystems, Inc.
STREET: 10301 Stella Link Rd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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100.0%; Score 77; DB 1; Length 15;
Best Local Similarity 100.0%; Pred. No. 8.5e-06;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                    COMPUTER: USA

ZIP: 22046-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/095,332
FILING DATE: 23-JUL-1993
CLLASSIFICATION: 435
FILING DATE: 13-JUL-1993
CLLASSIFICATION: 435
FRICA APPLICATION NUMBER: 07/760,530
FILING DATE: 18-SEP-1991
ATTORNEY/AGENT INFORMATION:
NAME: SVENBEND, Leonard R.
REGISTRATION NUMBER: 30,330
REFERENCE/POCKET UNBER: 1173-354p
TELECOMMUNICATION INFORMATION:
TELEFAK: 703-241-1300
TELEFEKX: 703-241-2848
STREET: Birch, Stewart, Kolash & Birch STREET: 301 N. Washington CITY: Falls Church STATE: Virginia COUNTRY: USA ZIP: 7.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 77025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7, Application US/08707801A Patent No. 5728814
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 248345
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: HIV-1
INDIVIDUAL ISOLATE: IIIB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 amino acide
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIQRGPGRAFVTIGK 15
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
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US-08-707-801A-7
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COUNTRY:
COUNTRY:
COUNTRY:
COMPUTER: BADABLE FORM:
MEDIUM TYPE: FLOPEY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,252
FILING DATE: O'LON-1935
APPLICATION NUMBER: 08/326,676
FILING DATE: 09-Jun-1995
APPLICATION NUMBER: 09/726,605
FILING DATE: 09-July-1991
APPLICATION NUMBER: 07/126,505
FILING DATE: 01-Mar-1991
APPLICATION NUMBER: 07/155,321
FILING DATE: 12-Peb-198
ATTOMREY/AGENT INFORMATION:
NAME: MAT'A C. H. Lin
REGISTRATION NUMBER: 29,323
REBERBURGE/DOCKET NUMBER: 1151-4004 US4
TELEPRANCE (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
TITLE OF INVENTION: OF USING SAME
TITLE OF INVENTION: HUMAN IMMUNOD
TITLE OF INVENTION: PROTEIN, DIAG
TITLE OF INVENTION: AND AS VACCIN
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
STREET: 345 PARK AVE.
CITY: NEW YORK
STATE: NEW YORK
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 2, Application US/08021879
; Patent No. 5817767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (212) 751-6845
TELEX: 421792
INPORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 15; Conservative
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JIY: New York
STATE: New York
COUNTRY: USA
ZIP: 1012
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US-08-488-252-27
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APPLICANT: Fung, Michael S.C., Sun, Bill N.C, Sun, Cecily R.Y.,
APPLICANT: Kin, Young W., Yu, Liming,
TITLE OF INVENTION: THE GCIQ RECEPTOR, HIV-1 gpl20 REGION BINDING
TITLE OF INVENTION: THERETO, AND RELATED PEPTIDES AND TARGETING
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Tanox Biosystems, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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Patent No. 5763160
GENERAL INFORMATION:
APPLICANT: Chang Yi Wang
ITILE OF INVENTION: SYNTHETIC PEPTIDES AND PROCESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 77; DB 1; I 100.0%; Pred. No. 8.5e-06;
                                                                           100.0%; Score 77; DB 1; I
larity 100.0%; Pred. No. 8.5e-06;
Conservative 0; Mismatcher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: TOWNSY: USA
ZIP: 77025
COUNTRY: USA
ZIP: 77025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch
COMPUTER: IBM PS/2
OPERATING SYSTEM: DOS 3.30
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/711,175
FILING DATE: 09-SEP-1996
CLASSIFICATION 5.30
PRIOR APPLICATION NUMBER: US/08/711,175
FILING DATE: 24-WAR-1995
ATTORNEY/AGENT INFORMATION:
PREFERENCE/DOCKET NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TNX95-1
TELECOMUNICATION INFORMATION:
TELEPHONE: (713) 664-2288
TELEPHONE: (713) 664-2288
TELEPHONE: (713) 664-8914
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Tanox Biosystems, Inc.
10301 Stella Link Rd.
                                                                                                                                                                                                                                                                                                                 ; Sequence 7, Application US/08711175
; Patent No. 5739306
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Matches 15; Conservative
               ; TYPE: amino acid
; TOPOLOGY: linear
US-08-709-006-7
                                                                                                                Best_Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 10
US-08-488-252-27
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                                                                                                                                                                                                                                                                                                         US-08-711-175-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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                                                                                                Query Match
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OF USING SAME FOR THE DETECTION OF ANTIBODIES TO HUMAN IMMUNODEFICIENCY VIRUS (HIV) GP120 ENVELOPE PROTEIN, DIAGNOSIS OF AIDS AND PRE-AIDS CONDITIONS AND AS VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gapa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Graham P. Allaway
APPLICANT: Graham P. Allaway
APPLICANT: Graham P. Allaway
TITLE OF INVENTION: SYNERGISTIC COMPOSITION OF CD4-BASED
TITLE OF INVENTION: PROTBIN AND ANTI-HIV-1 ANTIBODY, AND
TITLE OF INVENTION: METHODS OF USING SAME
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: ADDRESSEE: ADDRESSEE: ADDRESSEE: 30 Rockefeller Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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B: Tanox Biosystems, Inc.
10301 Stella Link Rd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 RIORGPGRAFVTIGK 15
                                 MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
ORGANISM: HIV-1
INDIVIDUAL ISOLATE: III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 713-664-8914 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                   LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                              NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: Linear
AMINO ACID
                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 10301
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Houston
STATE: Texas
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  77025
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US-07-950-571A-3
                                                                                                                                                                                 LOCATION:
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Sequence 1, Application US/07760530
Patent No. 5820865
APPLICANT: Berzofsky, Jay A. APPLICANT: Germain, Hidemi APPLICANT: Germain, Ronald N. TITLE OF INVENTION: SPECIFIC FOR A BROAD ARRAY OF HIV-1 ISOLATES USING HYBRID TITLE OF INVENTION: SPECIFIC FOR A BROAD ARRAY OF HIV-1 ISOLATES USING HYBRID TITLE OF INVENTION: SYNTHETIC PEPTIDES
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolash & Birch STRET: 301 N. Washington CITY: Falls Church STRET: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRATING TAPLICATION DATA:
APPLICATION NUMBER: US/07/760,530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 77; DB 2; I
100.0%; Pred. No. 8.5e-06;
tive 0; Mismatches 0;
              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/021,879
FILING DATE: 24-FEB-1993
ATTORNEY AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERRENCE/DOCKET NUMBER: 41189/JPW/AJM
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1173-354p
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ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30,330
REFERENCE/DOCKET NUMBER: 1173-
TELECPHONICATION:
TELECPHONE: 703-241-1300
TELEPHONE: 703-241-1300
                                                                                                                                                                                                                                                         TELEPHONE: (212) 977-9550
TELEPAX: (212) 664-0525
TELER: 42253 COPUI
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 248345
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.0
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 19910918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                             single
                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: protein US-08-021-879-2
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid STRANDEDNESS: 81
                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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US-07-760-530-1
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Fung, Michael S.C., Sun, Bill N.C., Sun, Cecily R.Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/07950571A

Batent No. S854400

GENERAL INFORMATION:
APPLICANT: Chang, Tse Wen, Fung, Michael S.C., Sun, Bill N.C., Sun, Cec. APPLICANT: Chang, Nancy T.
TITLE OF INVENTION: Monoclonal Antibodies which Neutralize HIV-1 Infection NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Tange Biografiens, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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/label= peptide
/note= "synchetic peptide, sequence = residues 315
/note= "synchetic peptide, sequence = residues 315
glycoprotein."
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                                                                                                                                                             Length 15,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Hi Density Diskette
COMPUTER: 1BM PS/2
COMPUTER: 1BM PS/2
COMPUTER: 1BM PS/2
COMPUTER: 1BM PS/2
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: No. 5854400 07/767,533
ATTORNEY AGENT INFORMATION:
NAME: MIYEABL, ELIC P.
REGISTRATION WHERE: 31,211
COMPUTED TO THE PROGRATION DATA:
NAME: MIXEABL, ELIC P.
REGISTRATION WHERE: 31,211
                                                                                                                                                       ch 100.0%; Score 77; DB 2; 1 Similarity 100.0%; Pred. No. 8.5e-06; 15; Conservative 0; Mismatches 0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 77; DB 2; Best Local Similarity 100.0%; Pred. No. 8.5e-06; Matches V.15; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TNX87-11BBC
TELECOMMUNICATION INFORMATION:
TELEPONE: 713-664-2288
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1 RIQRGPGRAFVTIGK 15

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Gaps
                                                                    APPLICANT: MATSUO, KAZUHIRO
APPLICANT: CHUJO, YOSHITOMO
APPLICANT: YAMAZAKI, AKIHIRO
APPLICANT: YAMAZAKI, SAUG
APPLICANT: YAMAZAKI, SHUDO
APPLICANT: YAMAZAKI, SHUDO
APPLICANT: TASAKA, HIROMICHI
TITLE OF INVENTION: ANTI-AIDS SECRETORY RECOMBINANT BCG
TITLE OF INVENTION: VACCINE
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON. SPIVAK, MCCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                                                                                                                                                                                                                                                                                                              STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400 CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/975,699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/JP95/01515
PRIOR DATE: 11-JUL-1995
PRIOR APPLICATION DATA: APPLICATION NUMBER: JP 178462/1994
PILING DATE: 29-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
REGISTATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-795-0X PCTELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION: 424
FIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/619,512
FILING DATE: 29-MAR-1996
APPLICATION NUMBER: PCT/JP95/01515
                   Sequence 6, Application US/08975699
Patent No. 5858369
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 15
US-08-972-089-6
; Sequence 6, Application US/08972089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 703-413-3000
TELEFAX: 703-413-2220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN: HIV-1 (JAPAN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM:
SOFTWARE: Patentl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
US-08-975-699-6
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APPLICANT: YAMAZAKI, AKIHIRO
APPLICANT: HONDA, MITSUO
APPLICANT: YAMAKAZI, SHUDO
APPLICANT: TASAKA, HIROMICHI
TITLE OF INVENTION: VACCINE
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSED: OBLON. SPIVAK, MCCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 8.5e-06;
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                                                                                                                                                                                                                  STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400 CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/972,089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRILING DATE:

CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/975,699
FILING DATE:
APPLICATION NUMBER: PCT/JP95/01515
FILING DATE: 31-JUL-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 178462/1994
FILING DATE: 29-JUL-1994
ATTONNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-795-0X PCT
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: peptide
ONGIGINAL SOURCE:
ORGANISM: HUMAN IMMUNODEFICIENCY VIRUS
STRAIN: HIV-1 (JAPAN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 7, Application US/08455625

Patent No. 593218

GENERAL INFORMATION:

APPLICANT: Berzofsky, Jay A.

APPLICANT: Pendlers, Jeffrey D.

APPLICANT: Pendlecon, C. D.

APPLICANT: Nara, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEBAX: 703-410 NO: SEQUENCE CHARACTERISTICS: FWATH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                                                                                                                                                    22202
                                                                                                                                                                                                                                                          STATE: VI
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us-08-869-386-1.rai

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TELEFAX: 212-596-9090 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , MOLECULE TYPE: peptide US-08-395-204-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: ami:
TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -08-628-687-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Pall Church
STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: 1.15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18IIIB peptide, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 77; DB 2; Length 15; Best Local Similarity 100.0%; Pred. No. 8.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner
STREET: 1300 I Street, N.W.
STREET: Washington
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Ladant, Daniel
APPLICANT: Leclerc, Claude
APPLICANT: Sebo, Peter
APPLICANT: Ullmann, Agnes
TITLE OF INVENTION: Recombinant Mutants for Inducing
TITLE OF INVENTION: Specific Immune Responses
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
TELEFAX: 703-205-8050
                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 17
US-08-395-204-2
; Sequence 2, Application US/08395204
; Patent No. 5935580
; Patent No. 5935580
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15 amino acida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide PRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-455-625-7
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Sequence 1, Application US/08628687
Sequence 1. Application US/08628687
Patent No. 5939277
GENERAL INFORMATION:
APPLICANT: Rakowing STATION:
TITLE OF INVENTION: DETECTION AND TREATMENT OF BREAST AND TITLE OF INVENTION: GYNECOLOGICAL CANCER NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS: ADDRESSEE: FISH & NEAVE STREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC compatible
COMPOTER: IBM PC compatible
COMPOTER: IBM PC compatible
COMPOTER: IBM PC compatible
CONTRARE: PatentIN Release #1.0, Version #1.25
SOFTWARE: PAPLICATION DATA:
APPLICATION NUMBER: US/08/395,204
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US 07/871,795
PRICK APPLICATION NUMBER: US 07/871,795
ATTORNEY/AGENT INFORMATION:
NAMBE: MCGENE, Kenneth J.
NAMBE: MCGENER, Kenneth J.
NAMBE: WCGENER, Kenneth J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SIREE: New York
STATE: New York
COUNTRY: USA
ZIP: 10020
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: PLOPSY disk
COMPUTER: PLOPSY disk
COMPUTER: PREPAIR PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/628,687
FILING DATE: 14-JUN-1996
CLASSIFICATION ADATA:
APPLICATION NUMBER: US 08/138,141
FILING DATE: 15-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: Haley Jr., James F.
REGISTRATION NUMBER: 27,794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 77; DB 2; Best Local Similarity 100.0%; Pred. No. 8.5e-06 Matches 15; Conservative 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                         NAME: Meyers, Kenneth J.
REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET UNBER: 03495-0109-00000
TELECOMMUNICATION: 1NFORMATION:
TELEPHONE: 202-408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: APPOLLO/1CIP1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
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peptide of HIV-I envelope glycoprotein from strain IIIB; activatible by protease cleavage to core
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GENERAL INFORMATION:
APPLICANT: Wallen:
TITLE OF INVENTION: Methods for Synthesizing Heat Shock Protein Complexes FILE REFERENCE: UNME-0008-1
CURRENT PILION NUMBER: US/08/986,234
CURRENT PILION DATE: 1997-12-05
NUMBER OF SEQ ID NOS: 114
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 13
LENGTH: 15
                                                                                                       /label= peptide
//note= "Highly immunogenic core peptide from
immunodominant region of envelope glycoprotein of
HIV-1 strain IIIb; peptide p18-1-10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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100.0%; Pred. No. 8.5e-06;
tive 0; Mismatches 0;
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100.0%; Pred. No. 8.5e-06;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label= peptide
/note= "peptide p18-V-10'
                                                                                                                                                                                                                                                         /label= peptide
/note= "peptide p18-I-9"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= peptide
/note= "peptide p18-V-9"
                                                                                                                                                                                                                                                                                                                                                                 /label= peptide
/note= "peptide p18-T-9"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ; ORGANISM: Human immunodeficiency virus US-08-986-234-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 28, Application US/08986234 Patent No. 5981706 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAWE/KEY: Peptide
LOCATION: 2..11
OTHER INFORMATION: /
OTHER INFORMATION: /
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                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: Peptide
LOCATION: 3.11
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                      LOCATION: 4.13
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
PEATURE:
                                                                                                                                                                                                               NAME/KEY: Peptide
LOCATION: 5.13
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                               LOCATION: 4..12
OTHER INFORMATION:
OTHER INFORMATION:
                        OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: Peptide
LOCATION: 4..12
                                                                  NAME/KEY: Peptide
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US-08-986-234-13
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APPLICANT: Berzofeky, Jay A.

APPLICANT: Takeshita, Toohiyuki

APPLICANT: Shirai, Muteunori

APPLICANT: Shirai, Muteunori

APPLICANT: Shirai, Muteunori

APPLICANT: Roalvowski, Stewen

APPLICANT: Margulies, David H.

TITLE OF INVENTION: Potent Peptide for Stimulation of

TITLE OF INVENTION: Cytotoxic T Lymphocytes Specific for the HIV-I Envelope

NUMBER OF SEQUENCES: 20

CORRESPONDENCE ADDRESS:

ADDRESSEE: Birch, Stewart, Kolash & Birch
                                                                                                                                                                                                                                                                                      ö
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/note= "Cytotoxic T lymphocyte immunodominant
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ZIP: 22046-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/847,311A
FILING DATE: 06-MAR-1992
CLASSIFICATION: 435
ATTONEY/AGENT INFORMATION:
NAME: SVENSEON, Leonard R.
REGISTRATION NUMBER: 30,330
REFERENCE/DOCKET NUMBER: 30,330
REFERENCE/DOCKET NUMBER: 1173-392P
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
word: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRAGMENT TYPE: internal ORIGINAL SOURCE: ORGANISM: Human Immunodeficiency Virus Type I
                                                                                                                                                                                                                                                                  . 8.5e-06;
                                                                                                                                                                                                                                            DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Birch, Stewart, Kolash & Birch STREET: 301 N. Washington CITY: Falls Church STATE: Virginia COUNTRY: USA
                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                          Query Match 100.0%; Score 77; Best Local Similarity 100.0%; Pred. No. Matches 15; Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/07847311A Patent No. 5976541
                                                                                                                                                                                                                                                                                                                                1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                     1 RIORGPGRAFVTIGK 15
                              LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                   ; ANTI-SENSE: NO; FRAGMENT TYPE: internal US-08-628-687-1
             SEQUENCE CHARACTERISTICS
                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: Peptide
LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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us-08-869-386-1.rai

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APPLICANT: Zaliphky, Samuel; Woodle, Martin, Martin, Francis;
APPLICANT: Zaliphky, Yecheakel
TITLE OF INVENTION: Enhanced Circulation Effector Composition and
TITLE OF INVENTION: Method
NUMBER OF SEGUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OURENTING SYSTEM:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,332
FILING DATE: 7-UN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/316,436
FILING DATE: 29-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/035,443
PRIOR APPLICATION NUMBER: US 08/035,443
FILING DATE: 29-ARR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Mohr, Judy M.
REGISTRATION NUMBER: 38,563
REFRENCE/COKET NUMBER: 5325-0115.31
TELECOMMUNICATION INFORMATION:
  PatentIn Release #1.0, .Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 77; DB 3; 100.0%; Pred. No. 8.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SSEE: Dehlinger & Associates
F: 350 Cambridge Avenue, Suite 250
Palo Alto
  SOFTWARE: Patentin Release #1.0, "CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/493,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-480-332-1; Sequence 1, Application US/08480332; Patent No. 6180134; GENERAL INFORMATION:
                                                                                CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Price, Robert L.
REGISTATION NUMBER: 22,685
REFERENCE/DOCKET NUMBER: 715-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-684-1111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                         TELEFAX: 703-684-1124
INFORMATION FOR SEQ ID NO: 25
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIORGPGRAFVTICK 15
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
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TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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STATE: CA
COUNTRY:
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TITLE OF INVENTION: Methods for Synthesizing Heat Shock Protein Complexes
'FILE REFERENCE: UNME-0008-1
CURRENT APPLICATION NUMBER: US/08/986,234
CURRENT FILING DATE: 1997-12-05
NUMBER OF SEQ ID NOS: 114
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 28
                                                                                                                                                                                                                                                                                                                           Gaps
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Sequence 22, Application US/08492076A

Sequence 22, Application US/08492076A

GENERAL INFORMATION:

APPLICANT: Adman, Sally E.

APPLICANT: Richardson, Simon M.

TITLE OF INVENTION: No. 6060064el Proteinaceous Particles
FILE REFERENCE: 10180.60968

CURRENT APPLICATION NUMBER: US/08/492,076A

CURRENT PILION DATE: 1993-02656

BARLIER PILICATION NUMBER: PCT/GB93/02656

BARLIER PILION DATE: 1993-12-24

NUMBER OF SEQ ID NOS: 23

SOFTWARE: PatentIN Ver. 2.0

SEQ ID NO 22

LENGTH: 15
                                                                                                                                                                                                                                                                     Score 77; DB 2; Length 15;
Pred. No. 8.5e-06;
Mismatches 0; Indels
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100.0%; Pred. No. 8.5e-06;
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Patent No. 6127149
GENERAL INFORMATION:
APPLICANT: Koshida, Shogo
APPLICANT: Koshida, Shogo
ITLE OF INVENTION:
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: LOWE, PRICE, LEBLANC & BECKER
STREET: 99 CANAL CENTER PLAZA, SUITE 300
CITY: ALEXANDRIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , ORGANISM: Human immunodeficiency virus type 1
US-08-492-076-22
                                                                                                                                                                                  TYPE: PRT
; ORGANISM: Human immunodeficiency virus
US-08-986-234-28
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WIBJUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.0%;
Matches 15; Conservative 0
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Best Local Similarity 100.
Matches 15; Conservative
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USA
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Gaps

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Length 15; Indels

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APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Nara, Peter
APPLICANT: Nara, Peter
APPLICANT: Nara, Peter
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                        100.0%; Score 77; DB 3; Length 15; 100.0%; Pred. No. 8.5e-06;
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                                                                                   Indels
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100.0%; Pred. No. 8.5e-06;
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MEDIUM TYPE: DESKETCE
COMPUTER: BM COMPACTIBLE
COMPUTER: BM COMPACTIBLE
COMPUTER: BM COMPACTIBLE
COMPUTER: MINIOWSSS
SOFTWARE: FASTENCY MINIOWSSS
SOFTWARE: FASTENCY MINIOWSSS
SOFTWARE: FASTENCY MINIOWSSS
SOFTWARE: FASTENCY MINIORSR: US/08/060,988A
FILING DATE: US/08/060,988A
FILING DATE: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 28-JAN-1988
FILING DATE: 28-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Beattie, Ingrid A.
REGISTRATION VINBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022001
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                Sequence 7, Application US/08060988A Patent No. 6294322 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
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617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
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Best Local Similarity 100.
Matches 15, Conservative
                                                                                     Conservative
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                                          Query Match
Best Local Similarity
Matches 15; Conserva
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US-08-455-685-7
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Patent No. 6214347

GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A. APPLICANT: Berzofsky, Jay A. APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                           Query Match 100.0%; Score 77; DB 3; Length 15; Best Local Similarity 100.0%; Pred. No. 8.5e-06; Matches 15; Conservative 0; Mismatches 0; Indels
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MEDIUM TYPE: Disketer
COMPUTER: IDB Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 13-MAY-1995
PRIOR APPLICATION NUMBER: 08/060,988
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/119,98
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 29-AUG-1998
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022003
TELECOMMUNICATION INFORMATION:
                                                                                          ANTI-SENSE: NO
ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: Peptide 1, Fig. 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Fish & Richardson P.C.
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                          1 RIQRGPGRAFVTIGK 15
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617/542-8906
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TYPE: amino acid
TOPOLOGY: linear
       STRANDEDNESS: unknown
                          TOPOLOGY: unknown
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                   ;
NAME/KEY: CDS
;
LOCATION: 1..15
US-08-480-332-1
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STREET: L...
TTV: BOSTON
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US-08-455-685-7
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; TYPE: PRT
; ORGANISM: HIV p18
US-09-827-688-9
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GENERAL INCRMATION:
GENERAL INCRMATION:
GENERAL INCRMATION:
GENERAL INCRMATION:
GENERAL INCRMATION:
TITLE OF INVENTION:
FILE REPERENCE:
CURRENT APPLICATION NUMBER: US/09/389,390
CURRENT FILING DATE: 1999-09-03
FRIOR APPLICATION NUMBER: 60/099,261
FRIOR PILING DATE: 1999-06-10
FRIOR FILING DATE: 1999-06-10
FRIOR FILING DATE: 1999-06-10
SOFTWARE: PALENTIN NUMBER: 60/139,045
FRIOR FILING DATE: 1999-06-10
SOFTWARE: PALENTIN VEY: 2.1
SEQ ID NO 1
LENGTH: 15
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                                                                                                                                                                               APPLICANT: Kim, Tae-Youn
APPLICANT: Chee, Ki-Young
APPLICANT: Chee, Ki-Young
APPLICANT: Cheng, Jin-Soo
APPLICANT: Chois, Sung-Yoo
APPLICANT: Chois, Wasong
APPLICANT: Chois, Wasong
APPLICANT: Chois, Mang-Seok
APPLICANT: Chois, Hong-Seok
APPLICANT: Chois, Hong-Seok
APPLICANT: Chois, Hong-Seok
APPLICANT: Chois, Wasong
APPL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 77; DB 3; Length 15;
Pred. No. 8.5e-06;
; Mismatches 0; Indels
                                                                                                                                  Mogam Biotechnology Research Institute
Kim, Tae-Youn
Lee, Ki-Young
Chang, Jin-Soo
Cho, Sung-Yoo
Hwang, Yu-Kyeong
Choi, Myeong
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US-09-508-552-15
'Sequence 15, Application US/09508552
'Patent No. 6749856
                            Sequence 8, Application US/09051006
Patent No. 6380359
GENERAL INFORMATION:
APPLICANT: Mogam Biotechnology Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 1, Application US/09389390
; Patent No. 6558961
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Best Local Similarity 100.0%;
Matches 15; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: PRT
; ORGANISM: HIV
US-09-051-006-8
US-09-051-006-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-389-390-1
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VS-09-827-688-9

Sequence 9, Application US/09827688

Sequence 9, Application US/09827688

Patent No. 6821955

GENERAL INFORMATION:
APPLICANT: KINSEY, BERMA

APPLICANT: KINSEY, BERMA

TITLE OF INVENTION: MACROAGGREGATED PROTEIN CONJUGATES AS ORAL GENETIC IMMUNIZATION DE

TITLE OF INVENTION: AGENTS

FILE REFERENCE: PO194949151/10004014

CURRENT PELING DATE: 2001-04-06

PRIOR APPLICATION NUMBER: 60/195,680

PRIOR FILING DATE: 2000-04-07

NUMBER OF SEQ ID NOS: 13

SOFTWARE: Patentin version 3.0

SOFTWARE: Patentin version 3.0

SOFTWARE: Patentin version 3.0
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APPLICANT: Berzofsky, Jay A.
APPLICANT: Belyakov, Igor M.
APPLICANT: Belyakov, Igor M.
APPLICANT: Derby, Michael A.
APPLICANT: Derby, Michael A.
APPLICANT: Strober, Warren
APPLICANT: THE GOVERNMENT OF THE UNITED STATES OF AMERICA, as
TITLE OF INVENTY BOUGHNERT OF THE UNITED STATES OF AMERICA, as
TITLE OF INVENTY BOUCOSAL CYTOTOXIC T LYMPHOCYTE RESPONSES
FILE REFERENCE: 36200PCSEQ
CURRENT FILING DATE: 1900-06-12
PRIOR APPLICATION NUMBER: 60/058,523
PRIOR APPLICATION NUMBER: 60/058,523
PRIOR APPLICATION NUMBER: 60/074,894
PRIOR FILING DATE: 1998-02-17
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 15
LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: LENGTH: 
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100.0%; Score 77; DB 4; I
Best Local Similarity 100.0%; Pred. No. 8.5e-06;
Matches 15; Conservative 0; Mismatches 0;
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100.0%; Pred. No. 8.5e-06;
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GENERAL INFORMATION:
APPLICANT: BOARD OF REGENTS, THE UNIVERSITY OF
APPLICANT: TEXAS SYSTEM
APPLICANT: AARLINGHAUS, Ralph B.
APPLICANT: ARLINGHAUS, Ralph B.
APPLICANT: ALINGHAUS, Chris D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT
; ORGANISM: Human immunodeficiency virus type 1
US-09-508-552-15
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Best Local Similarity 100.0
....hes 15; Conservative
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APPLICANT: Krafft, G.A.
APPLICANT: Mandecki, W.
APPLICANT: March, S.C.
APPLICANT: Rusell, J.R.
APPLICANT: Yue, V.T.
APPLICANT: Yue, V.T.
TITLE OF INVENTION: Genetically Engineered Enzymes And Their
TITLE OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 1.15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18IIIB peptide, see Table V"
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100.0%; Pred. No. 8.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                   APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
CLASSIPICATION: PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY AGENT INFORMATION:
NAME: SVENBSON, LEONBACK R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Softpo:
SOFTWARE: Softpo:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/657,392
FILING DATE:
FILING DATE:
PRIOR APPLICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/100,708
FILING DATE: July 29, 1993
                                                                                                                                                                                                                                 NAME: Svenseon, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
TELEPKX: 703-205-8050
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COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: SOftPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 28, Application US/08657392
Patent No. 5843634
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
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Brennan, C.A.
Bridon, D.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 100.
Matches 15; Conservative
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
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STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
TOPOLOGY: linear
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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Sequence 7, Application PC/TUS9405142
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSE: ADDRESSE:
TITLE OF INVENTION: STEWART, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STREET: Virginia
                                TITLE OF INVENTION: METHODS AND COMPOSITIONS
TITLE OF INVENTION: FOR ELICITING IMMUNE OR ANTI-INFECTIVE RESPONSES
TOWNER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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100.0%; Pred. No. 8.5e-06
tive 0; Mismatches 0
                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10378
FILING DATE: 19921202
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/800,932
FILING DATE: December 2, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/945865
FILING DATE: September 16, 1992
ATTORNEY AGENT INFORMATION:
NAME: PAFKEY, DAVIG L.
REGISTRATION NUMBER: 32,165
REGISTRATION NUMBER: 32,165
REGISTRATION INFORMATION:
TELECOMMUNICATION 
TELECOMMUNICATION
                                                                                                                                                         ADDRESSEE: Arnold, White & Durkee
STREET: P.O. Box 4433
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 22040-0747
COMPUTER READABLE FORM: ...
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0.
   NEHETE, Pramod N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVTIGK 15
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                       STATE: Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCT-US92-10378-1
                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Gaps

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Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1300 I Street, N.W., Suite 700 CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 35:
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                                                                                                  Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
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LENGTH: 16 amino acids
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                                           ENV 312-327
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Best Local Similarity
Matches 15; Conserv
      HIV-1
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      ORGANISM:
                    FEATURE:
;
LOCATION:
US-08-251-472-2
                                                                                                                                                                                                                                                                               RESULT 35
US-08-484-905-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-484-905-35
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                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: BUTILLON, CHRISTOPHE, MARTINON,
APPLICANT: FREDERIC; GRAS-MASSE, HELENE;
APPLICANT: GOMARD, ELISABETH, SERGHERAERT,
APPLICANT: CHRISTIAN; MAGNE, REMY; TARTAR,
APPLICANT: ANDRE; LEVY, JEAN-PAUL
TITLE OF INVENTION: CYTOTOXIC T LYMPHOCYTE
TITLE OF INVENTION: -INDUCING LIPOPEPTIDES AND USE AS VACCINES
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 2; Length 16; 100.0%; Pred. No. 9.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10016
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPEN DISK
COMPUTER: IBM FOR POSS MS-DOS
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/251,472
FILING DATE: 31-MAY-1994
CLASSIFICATION: 424
                                                               5324.US.P1
                                                                                                                                                        INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENCTH: 16 antho acid residues
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: BIERMAN & MUSERLIAN
STREET: 600 THIRD AVENUE
CITY: NEW YORK
STATE: NEW YORK
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/08251472
Patent No. 5871746
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: MUSERLIAN, CHARLES A
REGISTRATION NUMBER: 19,683
REFERENCE/DOCKET NUMBER: 102
TELECOMMUNICATION INFORMATION:
TELEFHONE: (212) 661-8000
TELEFAX: (212) 661-8002
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
ATTORNEY/AGENT INFORMATION:
NAME: Wong, Wean Khing
REGISTRATION NUMBER: 33.561
REFERENCE/DOCKET NUMBER: 5324
TELECHMUNICATION INFORMATION:
TELEPHONE: (708) 938-3517
TELEFFAX: (708) 938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 RIORGPGRAFVTIGK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
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Best Local Similarity 100.u
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                               TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                           US-08-657-392-28
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                                                                                                                                                                                                                                                                                                                     Sequence 35, Application US/08484905
Patent No. 5976551
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Abastado, Jean-Pierre
APPLICANT: Kourilsky, Philippe
TITLE OF INVENTION: An Altered Major Histocompatibility
TITLE OF INVENTION: Determinant
TITLE OF INVENTION: Determinant
NUMBER OF SEQUENCES: 127
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESSEE: Dunner
                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
100.0%; Score 77; DB 2; Length 16; 100.0%; Pred. No. 9.1e-06;
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                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Washington STATE: 0.005-315

ZIP: 2005-315

ZIP: 2006-315

COMPUTER EADABLE FORM:
MEDIUM TYPE: Floppy Disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS-MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 07/804,905
FILING DATE: 07-JUNE-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/801,818
FILING DATE: 05-DEC-1991
CLASSIFICATION: 530
PRIOR APPLICATION: 530
PRIOR APPLICATION S30
PRIOR APPLICATION: 530
PRIOR APPLICATION: 1830
FILING DATE: 15-NOV-1991
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: POCKET VUMBER: 33,332
REGISTRATION NUMBER: 33,332
REGISTRATION NUMBER: 33,332
REFERENCE/COMMUNDER: 03495.0106-03000
TELECOMMUNICATION: INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 2; I
100.0%; Pred. No. 9.1e-06;
tive 0; Mismatches 0;
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STATE: D.C.
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 RIQRGPGRAFVTIGK 15
                                     ADDRESSEE: BIERMAN & MUS
STREET: 600 THIRD AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 661-8002
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: ENV 312-327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   I: 16
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 15; Conserva
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                                                                                                                                          NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: HIV-1
                                                                                                            NEW YORK
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                                                                                                                                                                                                              10016
                                                                                                                                          STATE: NI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-248-082-2
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APPLICANT: BOUTLILON, CHRISTOPHE; MARTINON,

APPLICANT: BRIDERIC; GRAS-MASSE, HELENE;

APPLICANT: GONARD, ELISABETH; SERGHERAER;

APPLICANT: CHRISTIAN; MAGNE, REMY; TARTAR,

APPLICANT: ANDRE; LEVY, JEAN-PAUJ

TITLE OF INVENTION: CYTOTOVIC T LYMPHOCYTE

TITLE OF INVENTION: -INDUCING LIPOPEPTIDES AND USE AS VACCINES
                                                                                                                                                                                    APPLICANT: Mottez, Estelle
APPLICANT: Abastado, Jean-Pierre
APPLICANT: Kourilsky, Phillipe
TITLE OF INVENTION: Altered Major Histocompatibility Complex
TITLE OF SEQUENCES: 148
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 3; Length 16;
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Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: WABLILLINGLOID.

STATE: D.C.

ZIP: 20005-3315

COMPUTER READBABE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PATA:
APPLICATION NUMBER: US/08/481,985B
FILING DATE: 07-UN-1995
CLASSIFCATION: 435
PRIOR APPLICATION NUMBER: US 07/801,818
FILING DATE: 05-DEC-1991
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/801,818
FILING DATE: 05-DEC-1991
CLASSIFICATION DATA:
APPLICATION NUMBER: US 07/792,473
FILING DATE: 15-NOV-1991
CLASSIFICATION: 435
RIGHT APPLICATION NUMBER: US 07/792,473
FILING DATE: 15-NOV-1991
CLASSIFICATION: 435
APPLICATION: 435

                                                                                                                                                                                                                                                                                                                                                                                                                                           SSEE: Finnegan, Henderson, Farabow, Garrett
SSEE: Dunner
F: 1300 I Street, N.W., Suite 700
Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Meyers, Kenneth J.
REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 03495.0106-04000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%; Score 77; Best Local Similarity 100.0%; Pred. No. Matches 15; Conservative 0; Mismatch
                                                                                                 Sequence 35, Application US/08481985B
Patent No. 6011146
GENERAL INFORMATION:
APPLICANT: Mottez, Estelle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-248-082-2; Sequence 2, Application US/09248082; Patent No. 6015564
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  202-408-4400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                              JS-08-481-985B-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            셤
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APPLICANT: Abserted, Jean-Pierre
APPLICANT: Kourilsky, Phillipe
APPLICANT: Kourilsky, Phillipe
APPLICANT: Lone, Yu-Chun
APPLICANT: Casrouge, David
APPLICANT: Casrouge, Armanda
TITLE OF INVENTION: Altered Major Histocompatibility Complex
TITLE OF INVENTION: 127
CORRESPONDENCE ADDRESS:
ADDRESSER: Finnegan, Henderson, Farabow, Garrett &
ADDRESSER: Dunner
STREET: 1300 I Street, N.W., Suite 700
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 77; DB 3; 100.0%; Pred. No. 9.1e-06
                                                                                                                                                                                                  PILING DAIL.
CLASIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/251,472
FILING DATE: 31-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: MUSERLIAN, CHARLES A
REGISTRATION NUMBER: 19,683
REFERENCE/POCKET NUMBER: 102.1511
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/248,082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 6153408
GBNRAL INFORMATION
APPLICANT: Mottez, Estelle
APPLICANT: Abastado, Jean-Pierre
APPLICANT: Kourilsky, Phillipe
APPLICANT: Lone, Yu-Chun
APPLICANT: Lone, Yu-Chun
APPLICANT: Ojcius, David
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PCT-US94-02539-28
; Sequence 28, Application PC/TUS9402539
; GENERAL INFORMATION:
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                                                                                                  Brate, E.M.
Brennan, C.A.
Bridon, D.P.
Jaffe, K.D.
Krafft, G.A.
Mandecki, W.
March, S.C.
Russell, J.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: unknown
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                      APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCT-US94-02539-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATE: Terman, David S
TITLE OF INVENTION: SUPERANTIGEN BSED METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TRREATMENT OF INPECTIOUS DISEASE
TITLE OF INVENTION: TRREATMENT OF INPECTIOUS DISEASE
CURRENT APPLICATION NUMBER: US/08/992,877
CURRENT PILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/044,074
PRIOR FILING DATE: 1997-04-17
NUMBER OF SEQ ID NOS: 78
SOFTWARE: PATENTIN VET. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
100.0%; Score 77; DB 3; Length 16;
Best Local Similarity 100.0%; Pred. No. 9.1e-06;
Matches 15: Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9.1e-06;
                                                                                              CLASSIPICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 05 08/117,575
FILING DATE: 07-SEP-1933
APPLICATION NUMBER: US 08/072,787
FILING DATE: 06-JUN-1993
PRIOR APPLICATION DATA: APPLICATION NUMBER: US 07/801,818
FILING DATE: 05-DEC-1991
PRIOR APPLICATION NUMBER: US 07/792,473
APPLICATION NUMBER: US 07/792,473
ATTORNEY/AGENT INFORMATION:
NAME: MSYSTEM INFORMATION:
NAME: MSYSTEM INFORMATION:
NAME: MSYSTEM NUMBER: 25,146
REGESTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 052408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.0%; Score 77; DB 3; Best Local Similarity 100.0%; Pred. No. 9.1e-06 Matches 15; Conservative 0; Mismatches 0
                     SOFTWARE: Patentin Release #1.0, 1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/370,476
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15, Application US/08992877; Patent No. 6340461; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 35:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT . ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              , MOLECULE TYPE: peptide US-08-370-476-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-992-877-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ઠે
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Sequence 10. 5683695
GENERAL INFORMATION:
APPLICANT: Wang, Chang Yi
TITLE OF INVENTION:
TITLE OF SEQUENCES: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
TITLE OF INVENTION: Genetically Engineered Enzymes
TITLE OF INVENTION: And Their
TITLE OF INVENTION: Conjugates For Diagnostic Assays
TITLE OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
ADDRESSE: ABBOTT LABORATORIES
STREET: One Abbott Park Road
CITY: Abbott Park
STATE: 111inois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 77; DB 5; I
100.0%; Pred. No. 9.1e-06;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                            ZIP: 16064-3500

ZIP: 60064-3500

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
COMPUTER: SOFTEM: PC-DOS/MS-DOS
SOFTEME: SOFTEM
ZOFTEME: SOFTEM
APPLICATION NUMBER: PCT/US94/02539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5324.PC.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Wong, Wean Khing
REGISTRATION NUMBER: 33,561
REFRENCE/DOCKET NUMBER: 5324
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 938-3517
TELEPAX: (708) 938-2623
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1.1e-05;
FILING DATE: 01-FEB-1993
ATTORNEY/AGENT INFORMATION:
NAME: Adriano, Sarah B.
REGISTRATION NUMBER: 34,470
REFERENCE/DOCKET NUMBER: 34,470
TELECOMMUNICATION INFORMATION:
TELEPAN: 310-445-1140
TELEPAN: 310-445-1140
INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
LENGTH: 20 minno acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 10154
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
COMPUTER: ISM PC COMPATIBLE
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/326,676
FILING DATE: 07-Jun-1995
APPLICATION NUMBER: 07/26,605
FILING DATE: 09-July-1991
APPLICATION NUMBER: 07/663,262
FILING DATE: 01-Mar-1991
APPLICATION NUMBER: 07/155,321
APPLICATION NUMBER: 07/155,321
APPLICATION NUMBER: 07/155,321
APPLICATION NUMBER: 29,323
REGISTRATION NUMBER: 29,323
REFERENCE/DOCKET NUMBER: 1151-400
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 28, Application US/08488252 Patent No. 5763160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIQRGPGRAFVTIGK 15
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Best Local Similarity 100.
Matches 15; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 43
US-08-488-252-28
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Sequence 3, Application US/08121054C
Patent No. 5637481
GENERAL INFORMATION: Ledbetter, Jeffrey A. APPLICANT: Ledbetter, Jeffrey A. APPLICANT: Hayden, Martha S. APPLICANT: Linaley, Peter S. APPLICANT: Linaley, Peter S. APPLICANT: Fell, Perry TITLE OF INVENTION: Expression Vectors Encoding Bispecific TITLE OF INVENTION: Expression Proteins and Methods of Producing Biologically TITLE OF INVENTION: Active Bispecific Fusion Proteins in a Mammalian Cell NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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COUNTRY: USA

ZIP: 90025

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

CORRESTING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/121,054C

TING DATE: 13-SEP-1993

TING DATE: 13-SEP-1993
                                                                                                                                                                                          COMPUTER REALABLE FURVA:

MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC Compatible
COMPOTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.25
SOCTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/015,770B
FLING DATE: 10-FEB-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Wilson, M. Lisa
REFERENCE/DOCKET NUMBER: 34,045
REFERENCE/DOCKET NUMBER: 2002
TELECOMMUNICATION INFORMATION:
TELEFAX: (516)273-2828
TELEFAX: (516)273-177
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
TOWNEY: 18 Amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Merchant & Gould
STREET: 11150 Santa Monica Blvd., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                       ADDRESSEE: United Biomedical, Inc. STREET: 25 Davids Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 RIORGPGRAFVTICK 18
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Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-015-770B-4
                                                                                                                                                            ZIP: 11788
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
                                                                                         CITY: Hauppauge
                                                                                                                                          USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 42
US-08-121-054C-3
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GENERAL INFORMATION:

APPLICANT: Chang Yi Wang
TITLE OF INVENTION:
SYNTHETIC PEPTIDES AND PROCESS
TITLE OF INVENTION:
OF USING SAME FOR THE DETECTION OF ANTIBODIES TO
TITLE OF INVENTION: HUMAN IMMUNODEFICIENCY VIRUS (HIV) GP120 ENVELOPE
TITLE OF INVENTION: PROTEIN, DIAGNOSIS OF AIDS AND PRE-AIDS CONDITIONS
TITLE OF INVENTION: AND AS VACCINES
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN
STREET: 345 PARK AVE.
CITY: NEW YORK
STRIET: NEW YORK
COUNTRY: USA
                                                             Gaps
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100.0%; Score 77; DB 1; Length 20; 100.0%; Pred. No. 1.1e-05;
                                                             Indels
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Query Match 100.0%; Score 77; DB 3; Length 20; Best Local Similarity 100.0%; Pred. No. 1.1e-05; Matches 15; Conservative 0; Mismatches 0; Indels
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APPLICANT: Ledbetter, Jeffrey A.
APPLICANT: Ledbetter, Jeffrey A.
APPLICANT: Linaley, Peter S.
APPLICANT: Linaley, Peter S.
APPLICANT: Linaley, Peter S.
APPLICANT: Linaley, Peter S.
APPLICANT: Gilliland, Lisa K.
TITLE OF INVENTION: EXPRESSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
TITLE OF INVENTION: FUSION PROTEINS IN A MAMMALIAN CELL
TITLE OF INVENTION: FUSION PROTEINS IN A MAMMALIAN CELL
TITLE OF INVENTION: PUSION PROTEINS IN A MAMMALIAN CELL
CURRENT APPLICATION NUMBER: US/09/549,067A
CURRENT FILING DATE: 1995-10-05
PRIOR FILING DATE: 1995-10-05
PRIOR FILING DATE: 1993-09-13
PRIOR FILING DATE: 1993-09-01
PRIOR FILING DATE: 1993-09-01
PRIOR FILING DATE: 1993-09-01
PRIOR FILING DATE: 1993-02-01
PRIOR FILING DATE: 1993-02-01
PRIOR FILING DATE: 1993-02-01
PRIOR FILING DATE: 1993-01-22
PRIOR FILING DATE: 1993-01-22
PRIOR FILING DATE: 1993-01-22
PRIOR FILING DATE: 1993-01-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
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; ORGANISM: Homo sapiens
US-09-813-659-3
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                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3, Application US/08539436;
Patent No. 613292
GENERAL INFORMATION:
APPLICANT: Ledbetter, Jeffrey A.
APPLICANT: Gilliand, Lisa K.
APPLICANT: Linsley, Peter S.
APPLICANT: Linsley, Peter S.
APPLICANT: Bajorath, Jurgen
APPLICANT: Fell, Perry
TITLE OF INVENTION: Expression Vectors Encoding Bispecific
TITLE OF INVENTION: Active Bispecific Fusion Proteins in a Mammalian Cell
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
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                                                                                                                                                 100.0%; Score 77; DB 1; Length 20; 100.0%; Pred. No. 1.1e-05; ive 0; Mismatches 0; Indels
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COUNTRY: USA

ZIP: 90025

COUNTRY: USA

MEDIUM TYPE: Floppy disk

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/539,436

FILING DATE: 05-COT-1995

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/121,054

FILING DATE: 13-SEP-1993

AFFORMEY AGENT INFORMATION:

NAME: Adriano, Sarah B.

REFERENCE/DOCKET NUMBER: 30436.18US01

TELEPHONE: 310-445-1140

TELEFAX: 310-445-1140

TELEFAX: 310-445-1140

TELEFAX: 310-445-1140

TELEFAX: 310-445-1165:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Merchant & Gould
STREET: 11150 Santa Monica Blvd., Suite 400
                                                                                                                                                                                                                                                                                     6 RIQRGPGRAFVTIGK 20
                                                                                                                                                                                                                                                      1 RIORGPGRAFVTIGK 15
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LENGTH: 20 amino acids
TYPE: Amino acids
STRANDEDNESS:
TOPOLOGY: Unknown
                                                                                                                                                                           Best Local Similarity 100.0
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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                                                         ; STRANDEDNE;
; TOPOLOGY:
US-08-488-252-28
                                                                                                                                                                                                                                                                                                                                                                                RESULT 44
US-08-539-436-3
                                                                                                                                                      Query Match
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5 RIQRGPGRAFVTIGK 19

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Sequence 3, Application US/09813659
; Sequence 3, Application US/09813659
; Batent No. 6482919
; GENERAL INFORMATION:
; APPLICANT: Hayden, Martha S.
; APPLICANT: Linsley, Peter S.
; APPLICANT: Linsley, Peter S.
; APPLICANT: Fell, H. Petrix
; APPLICANT: Fell, H. Petrix
; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
; TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC
; TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC
; TITLE OF INVENTION: AND METHOD OF 90/9813, 659
; CURRENT PELING DATE: 2001-03-21
; PRIOR APPLICATION NUMBER: 09/549,067
; PRIOR FILING DATE: 1995-10-05
; PRIOR FILING DATE: 1995-10-05
; PRIOR FILING DATE: 1993-09-13
; FRIOR SED ID NOS: 32
; SOFTWARE: PARENTI ON ON SED ID NOS: 32
; SOFTWARE: PARENTI ON THE SED ID NOS: 32
; SOFTWARE: PARENTI ON THE SED ID NOS: 32
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Gaps ö Length 20; 0; Indels

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1.2e-05;
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330 University Avenue, 6th Floor
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Patent No. 5866320
                                                          Sequence 4, Application US/08453745A
Patent No. 5866137
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
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                                                                                                                                          APPLICANT: Rovinski, Benjamin
APPLICANT: Haynes, Joel
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nes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & MCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: linear
US-08-453-745A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-470-419-25
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Rovinski, Benjamin
APPLICANT: Haynes, Joel
APPLICANT: Haynes, Joel
APPLICANT: Cao, Shi Xian
APPLICANT: Klein, Michel H
TITLE OF INVENTION: Retrovirus-Like Particles Containing
TITLE OF INVENTION: Modified Envelope Glycoproteins
NUMBER OF SEQUENCES: 7
CONRESPONDENCE ADDRESS:
ADDRESSEE: 310 University Avenue, 6th Floor
                                                                                                                                                                                                                      100.0%; Score 77; DB 4; Length 20; 100.0%; Pred. No. 1.1e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.2e-05;
thes 0; Indels
                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: MSG LK/
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PACHICATION DATA:
APPLICATION NUMBER: US/08/452,503A
FILING DATE: 30-MAY-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/073,526
FILING DATE: 09-JAN-1993
CLASSIFICATION A35
FILING DATE: 09-JAN-1993
CLASSIFICATION NUMBER: US 08/073,526
FILING DATE: 09-JAN-1993
CLASSIFICATION NUMBER: US 08/073,526
FILING DATE: 09-JAN-1993
CLESSIFICATION NUMBER: US 08/073,526
FILING DATE: 09-JAN-1993
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTER.ESPICS:
LEMETAL OF A MISSIES
FILING DATE: 07-JAN-100 AS 05-1163
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PRIOR FILING DATE: 1991-06-27
NUMBER OF SEQ ID NOS: 35
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                      5 RIQRGPGRAFVTIGK 19
                                                                                                                                                                                                                                                                                                                        1 RIORGPGRAFVTIGK 15
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Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
; TOPOLOGY: linear
US-08-452-503A-4
                                                                                                                          TYPE: PRT
CORGANISM: Homo sapiens
US-09-549-067A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
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APPLICANT: CAO, Shi-Xian
APPLICANT: CAO, Shi-Xian
APPLICANT: CAO, Shi-Xian
APPLICANT: PROSSON, Roy
APPLICANT: FERSON, Roy
APPLICANT: KLEIN, Michel H
TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
TITLE OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
APPLICANT: Haynes, Joel
APPLICANT: Cao, Shi Xian
APPLICANT: Klein, Michel H
TITLE OF INVENTION: Retrovirus-Like Particles Containing
TITLE OF INVENTION: Modified Envelope Glycoproteins
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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STREET: 330 University Avenue, 6th Floor
CITY: Toronto
CITY: Toronto
CONTRY: Canada
ZIP: MGG IR7
COMPUTER EADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: DAY COMPATION
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURSENT APPLICATION DATA:
APPLICATION NUMBER: US/08/453,745A
FILING DATE: 09-MAY-1995
CLASSIFICATION ATA:
APPLICATION NUMBER: US/08/073,526
FILING DATE: 09-JAN-1993
CLASSIFICATION TINFORMATION:
NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,73
REPERRENCE/DOCKET NUMBER: 1038-445 MIS:a8
TELECOMMUNICATION HUMCHAILON:
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Mismatches
                                                                                                                                                             | IMMEDIATE SOURCE: CLONE: HIV-V3 acceptor peptide US-08-648-298-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25, Application US/08761828
Patent No. 5879925
GENERAL INFORMATION:
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TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 aming acids
                                                                                                                                                                                                                                                                                                                                            1 RIQRGPGRAFVTIGK 15
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                                                                                                   MOLECULE TYPE: peptide ORIGINAL SOURCE: ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . Query Match
Best Local Similarity 100.0
Matches 15; Conservative
                        21 amino acids
                                                                                                                                                                                                                                                                              Best Local Similarity 100.
Matches 15; Conservative
                                            TYPE: peptide
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
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STATE: Ontario
COUNTRY: Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
STREET: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 51
US-08-761-828-25
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Fatent No. 5871990
GENERAL INFORMATION:
APPLICANT: Henrik Clausen
TITLE OF INVENTION: UDP-N-acetyl-alpha-D-galactosamine:polypeptide
TITLE OF INVENTION: N-acetylgalactosaminyltransferase GalNAc-T3
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSE: Darby & Darby PC
STREET: 805 Third Avenue
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: NY
ZIP: 10022
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM PC compatible
OPERATING SYSTEM: DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/648,298
FILING DATE: 15-JUN-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 77; DB 2; Length 21; Best Local Similarity 100.0%; Pred. No. 1.2e-05; Matches 15; Conservative 0; Mismatches 0; Indels
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,419
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/290,105
FILING DATE: AUGUST 1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
RECISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-385 MIS:jb
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/ACENT INFORMATION:
NAME: Green, Reza
REGISTRATION NUMBER: 38,475
REFERENCE/DOCKET NUMBER: 4035/0B865
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212527700
TELEFAX: 2127536237
                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (416) 595-1155
TELEPAX: (416) 595-1163
INPORMATION POR SEQ ID NO: 25:
SEGURNCE CHARACTERISTICS:
LENGTH: 21 amino acids
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
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APPLICANT: ROVINSKI, Benjamin
APPLICANT: CAO, Shi-Xian
APPLICANT: CAO, Shi-Xian
APPLICANT: YAO, Fei-Long
APPLICANT: YAO, Fei-Long
APPLICANT: KLEIN, Michel H
ITILE OF INVENTION: ANTIGENICALLY-MARKED NON-INPECTIOUS RETROVIRUS-LIKE PARTICLES
CORRESPONDENCE ADDRESS: ö Score 77; DB 2; Length 21; Pred. No. 1.2e-05; Length 21; Indels 0; Indels COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
FILING DATE:
FILING DATE: 100.0%; Score 77; DB 2; I 100.0%; Pred. No. 1.2e-05; CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/290,105
FILING DATE: 15-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-655 MIS:jb
TELEPHONE: (416) 595-1155 E: Sim & McBurney 6TH Floor, 330 University Avenue

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US-08-776-949-25
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1 Sequence 4, Application US/08452520B

2 Sequence 4, Application US/08452520B

3 Sequence 4, Application US/08452520B

3 Patent No. 5912338 S840872

GENERAL INFORMATION:

4 APPLICANT: Rovinski, Benjamin

APPLICANT: Cao, Shi Xian

APPLICANT: Klein, Michel H

7 TITE OF INVENTION: Retrovirus-Like Particles Containing

TITLE OF INVENTION: Modified Envelope Glycoproteins

NUMBER OF SEQUENCES: 7

CORRESPONDENCE ADDRESS:

Annoncement Canada 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
100.0%; Score 77; DB 2; Length 21;
Best Local Similarity 100.0%; Pred. No. 1.2e-05;
Matches 15; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPESSE:
ADDRESSE:
ADDRESSE:
ADDRESSE:
ADDRESSE:
ADDRESSE:
Sim & McBurney
STREET: 330 University Avenue, 6th Floor
CITY: Torontc
CITY: Torontc
CITY: Torontc
COMPUTRY: Canada
ZIP: MGG IR7
COMPUTRY: Thory disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CUMRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/452,520B
FILING DATE: 30-MX-1995
CLASSIFICATION: 435
PRIOR APPLICATION: 435
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 24,973
ATTORNEY/AGENT INFORMATION:
ANAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
ATTORNEY/AGENT INFORMATION:
TELEPAN: (416) 595-1163
INFORMATION FOR SEC IN 00: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: Jananar
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US-08-290-105-25
Sequence 25, Application US/08290105
Patent No. 5955342
GENERAL INFORMATION:
APPLICANT: ROVINSKI, Benjamin
APPLICANT: CAO, Shi-Xian
APPLICANT: YAO, Fei-Long
APPLICANT: PERSSON, ROY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: KLEIN, Michel H
TITLE OF INVENTION: ANTIGEN
TITLE OF INVENTION: RETROVI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 RIQRGPGRAFVTIGK 15
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7 RIORGEGRAFVTIGK 21
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CORRESPONDER ADDRESS:
ADDRESSES: Sim & McBurney
FIRET: Suite 701, 30 University Avenue
CTITY: Toronto
CTITY: To
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Gaps
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APPLICANT: ROVISKI, Benjamin
APPLICANT: CAO, Shi-Xian
APPLICANT: CAO, Shi-Xian
APPLICANT: CAO, Shi-Xian
APPLICANT: PERSSON, ROY
APPLICANT: RIBIN, Michel H
TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sime & McBurney
STREET: 6th Floor, 330 University Avenue
CITY: Torontco
STRATE: Ontario
COUNTRY: Canada
ZIP: MGG 1R7
COMPUTER: READALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BEREIT Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/027,955
FILING DATE: 15-AUG-1994
ATTORNEY/BGENT INFORMATION:
REGISTRATION NUMBER: US/09/027,955
FILING DATE: 15-AUG-1994
ATTORNEY/BGENT INFORMATION:
REGISTRATION NUMBER: 1038-798 MIS:jb
TELERONMUNICATION NUMBER: 1038-798 MIS:jb
TELERONMUNICATION NUMBER: 1038-798 MIS:jb
TELERONMUNICATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-798
TELERONMUNICATION NUMBER: 1038-798
TELERONMUNICATION NUMBER: 1038-798
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TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
RETROVIRUS-LIKE PARTICLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
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Patent No. 6342228
GENERAL INFORMATION:
APPLICANT: ROVINSKI, Benjamin
CAO, Shi-Xian
YAO, Fei-Long
PERSSON, ROY
                                                                                                                                                           Sequence 25, Application US/09027955
Patent No. 6291157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIQRGPGRAFVTIGK 15
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Best Local Similarity 100..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                             US-09-027-955-25
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US-09-636-805-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | Sequence 25, Application US/08482810
| Patent No. 6080408
| GENERAL INFORMATION:
| APPLICANT: ROVINSKI, Benjamin
| APPLICANT: CAO, Shi-Xian
| APPLICANT: YAO, Fel-Long
| APPLICANT: WAO, Fel-Long
| APPLICANT: MICHEN, My CAO, Fel-Long
| APPLICANT: MICHEN, Michel H.
| TITLE OF INVENTION: NIFECTIOUS BY A PLURALITY OF MUTATIONS
| NUMBER OF SEQUENCES: 26
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,810
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
REFERENCE/DOCKET NUMBER: 1038-673 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO. 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRICE APPLICATION DATA:
APPLICATION NUMBER: US 08/292,967
FILING DATE: 22-AUG-1994
CLASSIFICATION: 424
ATTORNEY AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTATION NUMBER: 24,973
REFERENCE DOCKET NUMBER: 1038-490 MIS: vg
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Sim & McBurney
Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELBFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
                                                                                                                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                  RIORGPGRAFVTIGK 21
                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: Suite 701, 330
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                            ropology:
                                                                                                                                                                                                                                    US-08-776-949-25
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TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS RETROVIRUS-LIKE PARTICLES
                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 77; DB 4; Length 21; 100.0%; Pred. No. 1.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-1065 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPRONE: (416) 595-1163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
PRIOR APPLICATION DATA:
APPLICATION WOMBER:
FILING DATE: 22-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-924 MIS:jb
TELEPONE: (416) 595-1155
TELEPAX: (416) 595-1165
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MONDUTER: IBM PC compartible
COMPUTER: IBM PC compartible
SOFRWARE: PatentIn Release #1.0, Ve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/027,955
FILING DATE: 23.FEB-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/635,754
FILING DATE: 10-Aug-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 59
US-09-632-754-25
US-09-632-754-25
Sequence 25, Application US/09635754
Patent No. 6518030
GENERAL INFORMATION:
APPLICANT: ROVINSKI, Benjamin
ZAO, Shi-Xian
YAO, Fei-Long
PERSSON, ROJ
PERSSON, ROJ
PERSSON, ROJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KLEIN, Michel H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 RIORGPGRAFVTIGK 21
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                            single
                                                                                                                                                                                                                                                                                                       amino acid
                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                       US-09-258-128-25
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Patent No. 6451322
BAPELICANT: ROVINSKI, Benjamin
APPLICANT: CAO, Shi-Xian
APPLICANT: TAO, Fei-Long
APPLICANT: PERSSON, ROY
APPLICANT: RIBIN, Michel H
TITLE OF INVENTION: INFECTIOUS BY A PLURALITY OF MUTATIONS
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                           COMPUTER: IEBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/636,805
FILING DATE: 10-Aug-2000
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION NUMBER: US 09/027,955
FILING DATE: 23-FEB-1998
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-1068 MIS:jb
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 21;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/258,128
FILING DATE:
         CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 77; DB 3; I
100.0%; Pred. No. 1.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 25:
                                                                          CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
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US-09-258-128-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PARTICLES MADE NON-
PLURALITY OF MUTATIONS
                                                                            Query Match 100.0%; Score 77; DB 4; Length 21; Best Local Similarity 100.0%; Pred. No. 1.2e-05; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 4; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
100.0%; Score 77; DB 4; Length 21
Best Local Similarity 100.0%; Pred. No. 1.2e-05;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: MSG IR7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/680,525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIPTICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UNDABR:
APPLICATION UNDABR: 108 08/292,967
FILING DATE: 22-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-617 MIS:jb
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
; SEQUENCE DESCRIPTION: SEQ ID NO: 25:
US-09-635-754-25
                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: ROVINSKI, Benjamin
APPLICANT: CAO, Shi-Xian
APPLICANT: VAO, Fei-Long
APPLICANT: KLEIN, Michel H
TITLE OF INVENTION: RETROVINUS-LIKE PAI
TITLE OF INVENTION: INFECTIOUS BY A PLIC
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 61
US-09-636-223-25
; Sequence 25, Application US/09636223
; Patent No. 6544752
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                 Sequence 25, Application US/08680525 Patent No. 6544527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25: SEQUENCE CHARACTERISTICS:
LENGTH: 21 aming acids
                                                                                                                                                                 1 RIORGPGRAFVTICK 15
                                                                                                                                                                                                          7 RIORGPGRAFVTIGK 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
ADDRESSEE: Sim & McI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TOPOLOGY: linear
US-08-680-525-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Ontario
COUNTRY: Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                 US-08-680-525-25
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Sequence 13, Application US/08125012
| Sequence 13, Application US/08125012
| Patent No. 5533972
| GENERAL INFORMATION:
| PAPLICANT: Weiner, David B. APPLICANT: Wainer, David B. APPLICANT: Wainer, David B. APPLICANT: Wang, Bin APPLICANT: Coney, Lealie R. ITTLE OF INVENTION: Genetic Immunization NUMBER OF SEQUENCES: 34
| CORRESPONDENCES: 34
| CORRESPONDENCES: Woodcock Washburn Kurtz Mackiewicz & No. 5593972ria STREET: One Liberty Place 46th Floor
| CITY: Philadelphia STATE: Pennsylvania COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ..
0
APPLICANT: ROVINSKI, Benjamin
CAO, Shi-Xian
YAO, Fei-Long
PERSON, ROY
KLEIN, Michel H
KLEIN, ANTIGENICALLY-MARKED NON-INFECTIOUS
TITLE OF INVENTION: ANTIGENICALLY-MARKED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Version #1.25
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APPLICATION NUMBER: US/09/636,223
FILING DATE: 29-Dec-2000
CLASSIFICATION DATA:

PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 09/027,955
ATTORNEY/AGENT INFORMATION:
NAME: STEWART: Michael I
REGISTRATION NUMBER: 24,973
REGISTRATION NUMBER: 24,973
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                   NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 77; DB 4; I 100.0%; Pred. No. 1.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
;
SEQUENCE DESCRIPTION: SEQ ID NO: 25:
US-09-636-223-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7 RIORGPGRAFVTIGK 21
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Matches 15; Conservative
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Sequence 13, Application US/08453349

Retent No. 5830876

GENERAL INFORMATION:
APPLICANT: Williams, William V.
APPLICANT: Williams, William V.
APPLICANT: Walliams, Williams, Will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: PLOPPy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/453,349
                                                                                                                REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: APOL-0013
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEFAX: 215-568-3429
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APOL-0013
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/029,336
APPLICATION NUMBER: 08/008,342
APPLICATION NUMBER: 08/008,342
FILING DATE: January 26, 1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Deluca, Mark
REGISTRATION NUMBER: 33,229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: ALTELECOMMUNICATION INFORMATION: TELEPHONE: 215-568-3100 TELEFAX: 215-568-3429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
       CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: DeLuca, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 215-568-3429
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-453-349-13
                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-783-818-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Philadelphia
STATE: Pennsylvania
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
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US-08-453-349-13
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APPLICANT: Weiner, David B.
APPLICANT: Williams, William V.
APPLICANT: Williams, William V.
APPLICANT: Williams, William V.
APPLICANT: Word, Bin
APPLICANT: Coney, Leslie R.
TITLE OF INVENTION: Genetic Immunization
NUMBER OF SEQUENCES:
ADDRESSE: Woodcock Washburn Kurtz Mackiewicz & No. 5817637ris
STREET: One Liberty Place 46th Floor
CITY: Philadelphia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: Permisyroning
CONDUTER: Permisyroning
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: Datentin Release #1.0, Version #1.25 mb-MD/JAF
SOFTWARE: Patentin Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/783,818
FILING DATE: 13-4AN-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/125,012
FILING DATE: 21-5EP-1993
APPLICATION NUMBER: 08/029,336
FILING DATE: 11-MAR-1993
CLASSIFICATION: 514
PRICR APPLICATION NUMBER: 08/029,336
FILING DATE: 11-MAR-1993
CLASSIFICATION NUMBER: 08/029,336
FILING DATE: 11-MAR-1993
CLASSIFICATION NUMBER: 08/029,336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                  SOFTWARE: Patentin Release #1.0, Version #1.25 mb-MD/JAF CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/125,012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 77; DB 1; Length 22; 100.0%; Pred. No. 1.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: APOL-0013
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEPKX: 215-568-3100
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13, Application US/08783818
Patent No. 5817637
                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/008,342
FILING DATE: 26-JAN-1993
                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/029,336
FILING DATE: 11-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100..
Best Local 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear;
; MOLECULE TYPE: peptide
US-08-125-012-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 63
US-08-783-818-13
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FILING DATE: 26-JAN-1993
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                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "This sequence corresponds
to 303 to 324 of gp120 from the IIIB isolate."
                                           .
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    Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ch 100.0%; Score 77; DB 2; Length 22; Similarity 100.0%; Pred. No. 1.2e-05; 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                     APPLICANT: ZOLLA-PAZNER, SUBBAN
APPLICANT: ZOLLA-PAZNER, SUBBAN
TITLE OF INVENTION: HETEROHYBRIDOMAS PRODUCING HUMAN
TITLE OF INVENTION: MONOCLONAL ANTIBODIES TO HIV-1
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Parentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/345,321
100.0%; Score 77; DB 2; L 100.0%; Pred. No. 1.2e-05; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Browdy and Neimark
STREET: 419 Seventh Street, N.W., Suite 300
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/872,675
FILING DATE:
ATTORNEY/AGBRT INPORMATION:
NAME: Browdy, ROGET L.
REGISTRATION NUMBER: 25,618
REFERENCE/POCKET NUMBER: 20.61A-PAZNERIB
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEPHONE: 202-737-3528
                                                                                                                                                                                                          US-08-345-321-2
; Sequence 2, Application US/08345321
; Patent No. 5914109
; GENERAL INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMpatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 202-737-3528
TELEX: 248633
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                   1 RIQRGPGRAFVTIGK 15
                                                                                                                          8 RIORGPGRAFVTICK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
INDIVIDUAL ISOLATE: II
                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: 1.22
CTHER INFORMATION:
CTHER INFORMATION:
US-08-345-321-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid TOPOLOGY: linear
                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20004
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  Query Match
                                           Matches
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RESULT 66

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Gaps
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STREET: One Liberty Place 46th Floor
CITY: Philadelphia
STATE: Pennsylvania
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/979,385B
FILING DATE: 26-NOV-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/495,684
FILING DATE: 28-88P-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/495,684
FILING DATE: 26-78P-1995
FILING DATE: 26-78P-1995
FILING DATE: 21-58P-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/033,235
FILING DATE: 15-401-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/033,235
FILING DATE: 15-401-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/033,235
FILING DATE: 11-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/033,235
FILING DATE: 11-MAR-1993
PRIOR APPLICATION NUMBER: 08/029,336
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Sequence 11, Application US/08979385B
Patent No. 5981505
GENERAL INFORMATION:
APPLICANT: Weiner, David B.
APPLICANT: Williams, William V.
APPLICANT: Wang, Bin
TITLE OF INVENTION: Compositions and Methods for Delivery of
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 26-JAN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 1, Application US/08537245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: UPAP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEFAX: 215-568-3429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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SEQUENCE CHARACTERISTICS:
FUNCTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                        19103
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US-08-537-245-1
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APPLICATION NUMBER: US/08/805,889
                      FILING DATE: 03-MAR-1997
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,314
FILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: VENEKO, NANCY WAYS
REGISTRATION NUMBER: 36,298
REFERENCE/DOCKET NUMBER: NIHO
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-235-8556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 22 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 619-235-0176
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
FRAGMENT TYPE: internal
US-08-805-889-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92660
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APPLICANT: Earl, Patricia L.
APPLICANT: Barl, Patricia L.
APPLICANT: Dome, Robert W.
TITLE OF INVENTION: Oligomeric HIV-1 Envelope Glycoproteins
NUMBER OF SEQUENCES:
OCRRESPONDENCE ADDRESS:
ADDRESSEE: Knobbe, Martens, Olson and Bear
STREET: 620 Newport Center Drive 16th Floor
CITY: Newport Beach
STATE: CA
STATE: Pacentin Replay disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BMP PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 77; DB 2; Length 22; 100.0%; Pred. No. 1.2e-05; tive 0; Mismatches 0; Indels
  BATENT NO. 5985275
GENERAL INFORMATION:
APPLICANT: Neurath, A. Robert, Debnath, Asim K.,
APPLICANT: Jiang, Shibo
TITLE OF INVENTION: Proteins and Peptides Modified By
TITLE OF INVENTION: Aromatic Acid Anhydride Compounds
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Frishauf, Holtz, Goodman & Woodward
STREET: 600 Third Avenue
                                                                                                                                                                                                                                                                                                     COUNTY 10016
ZIP 10016
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3+ inch, 0.72 mb storage
COMPUTER: IBM PC
COMPUTER: IBM PC
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/420,573
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: BAICH, Richard
REGISTRATION NUMBER: 28,180
REFERENCE/DOCKET NUMBER: 950157/RSB
TELECOMONICATION INFORMATION:
TELEPHONE: (212) 972-1400
                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC
OPERATING SYSTEM: MS DOS
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/537,245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: cDNA to genomic RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 5, Application US/08805889 Patent No. 6039957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 370-1622
TELEX: 236268
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                    SIRBEA. CITY: New York STATE: New York COMMIRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                             COUNTRY:
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Sequence 5, Application US/09070291

Patent No. 6171596

GENERAL INPORMATION:
APPLICANT: Bards, Patricia L.
APPLICANT: Broder, Christopher C.
APPLICANT: Moss, Robert W.
APPLICANT: Moss, Bernard
ITLE OF INVENTION: Oligomeric HIV-1 Envelope Glycoproteins
NUMBER OF SEQUENCES: 10
CORRESPONDERSER: ADDRESSER: ADDRESSER: ADDRESSER: G20 Newport Center Drive 16th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 77; DB 3; I
100.0%; Pred. No. 1.2e-05;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AUDRESSEE: Knobbe, Martens, Olson and Bear STREET: 620 Newport Center Drive 16th Floor CITY: Newport Beach STATE: CA
NAME: FULLET, MICHAEL L.
REGISTRATION NUMBER: 36,516
REFERENCE/DOCKET NUMBER: 36,516
REFERENCE/DOCKET NUMBER: NIH079.001A
TELECOMMUNICATION INFORMATION:
TELEPAR: 619-235-8550
INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acids
TYPE: amino acids
STRANDEDNESS: single
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; Sequence 1, Application US/08097751; Patent No. 5527666; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Costigan, James V. REGISTRATION NUMBER: 25,669
REFERENCE/DOCKET NUMBER: 519
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 RIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide

US-08-880-576-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-097-751-1
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Sequence 13, Application US/08880576
Sequence 13, Application US/08880576
Sequence 13, Application US/08880576
GENERAL INFORMATION:
APPLICANT: Weiner, David B.
APPLICANT: Wang, Bin
APPLICANT: Wang, Bin
TITLE OF INVENTION: Genetic Immunization
TITLE OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 6468982ris
STREET: One Liberty Place 46th Floor
CITY: Philadelphia
STATE: Pennsylvania
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                        Sequence 22. Application US/09217306B
| Sequence 22. Application US/09217306B
| Patent NO. 6465220
| GENERAL INFORMATION:
| APPLICANT: Hassan, Helle
| APPLICANT: Clausen, Henrik
| APPLICANT: Bennet, Eric P.
| TITLE OF INVENTION: Glycosylation Using GalNAc-T4 Transferase
| FILE REFERENCE: 8850*1
| CURRENT APPLICANT: 1998-12-21
| NUMBER OF SEQ ID NOS: 25
| SOFTWARE: Patentin version 3.1
| SEQ ID NO 22
| LENGTH: 22
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYETE: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/880,576
FILING DATE: 23-UNN-1997
                                                                                                                                Query Match
100.0%; Score 77; DB 3; Length 22;
Best Local Similarity 100.0%; Pred. No. 1.2e-05;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 77; DB 4; Length 22;
Pred. No. 1.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) NAME/KEY: MISC FEATURE
; OTHER INFORMATION: HIVIIIB gp120
US-09-217-3068-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; So
Best Local Similarity 100.0%; Pi
Matches 15; Conservative 0;
                                                                                                                                                                                                                     1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                     8 RIQRGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 RIORGPGRAFVTIGK 17
                                           ANTI-SENSE: NO
FRAGMENT TYPE: internal
MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19103
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US-08-880-576-13
                                                                    ; FRAGMENT TY
US-09-070-291-5
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APPLICANT: Pasti, Marcella
APPLICANT: Mammano, Fabrizio
APPLICANT: Panozzo, Marina
APPLICANT: Panozzo, Marina
APPLICANT: Dettin, Monica
APPLICANT: Disello, Carlo
APPLICANT: Chieco-Bianchi, Luigi
ITILE OF INVENTION: METHOD FOR THE DIACNOSIS IN VITRO OF
TITLE OF INVENTION: HIV-1 VIRUS INFECTIONS
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSE: Hedman, Gibson, Costigan & Hoare
STREET: 1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
COMPUTER: IBM PS/2
OPERATING SYSTEM: DOS
SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 77; DB 4; I
100.0%; Pred. No. 1.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Hedman, Gibson, Costigan & Hoare STREET: 1185 Avenue of the Americas CITY: New York COUNTRY: USA YORK COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
CLASSIFICATION: 514
PRIOR APPLICATION DATA
PRIOR APPLICATION NUMBER: 08/125,012
FILLING DATE: 11-MAR-1993
APPLICATION NUMBER: 08/029,336
FILING DATE: 11-MAR-1993
PRIOR APPLICATION NUMBER: 08/008,342
FILING DATE: 26-JAN-1993
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Debluca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: APOL-0013
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3429
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/097,751
FILING DATE: 19930723
FILIASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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; TOPOLOGY:
US-08-257-528B-99
                  RESULT 74
US-08-257-528B-99
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                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Mastico, Robert Allan
APPLICANT: Stockley, Peter George
APPLICANT: Talbot, Simon John
TITLE OF INVENTION: Antigen-Presenting Capsid with
TITLE OF INVENTION: Fusion MS2-Coat Protein
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                            1.3e-05;
                                                                                                                                                                                                Query Match 100.0%; Score 77; DB 1; Best Local Similarity 100.0%; Pred. No. 1.3e-05 Matches 15; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: DISACETCE, 3.5", 1.44MD
COMPUTER: DISACETCE, 3.5", 1.44MD
COMPUTER: DEM PS2-486
COMPUTER: DEM PS2-486
COMPUTER: MS-DOS
SOFTWARE: WOORDERFECE 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/090,148
FILING DATE: 08/11/93
CLASSIFICATION 435
PRIOR APPLICATION A135
PRIOR APPLICATION NUMBER: GB 9101550.3
FILING DATE: 01/24/91
APPLICATION NUMBER: PCT/GB92/00124
FILING DATE: 01/24/92
ATTONEY/AGENT INPORMATION:
NAME: Nissenbaum, Israel
REGISTRATION NUMBER: 27,582
                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-090-148-6; Sequence 6, Application US/08090148; Patent No. 5534257; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Rosenman & Colin
STREET: 575 Madison Avenue
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 940-8636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (212) 940-6404 INFORMATION FOR SEQ ID NO: 6:
(212) 302-8989
                     TELEFAX: (212) 302-8998
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8 RIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                              1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                      8 RIORGPGRAFVTIGK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 24 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: NY
COUNTRY: U.S.A.
ZIP: 10022-2585
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , MOLECULE TYPE: PEPTIDE US-08-090-148-6
                                                                                                              ; TYPE: amino acid
; TOPOLOGY: circular
US-08-097-751-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: AMINO ACID
TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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Sequence 99, Application US/08460602A

Sequence 99, Application US/08460602A

PAPLICANT: S1A, Charles D.Y.

APPLICANT: CHONG, Pele

APPLICANT: CHONG, Pele

TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides

NUMBER OF SEQUENCES: 101

CORRESPONDENCE SUBCRES: 310

STREET: Suite 701, 330 University Avenue
                                      GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: CHONG, Pele
APPLICANT: CHONG, Pele
APPLICANT: TAIRIN Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FO Compatible
OPERATUR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/257,528B
FILING DATE: 09-UN-1994
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 1; I 100.0%; Pred. No. 1.3e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1038-336 MIS:jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Toronto STATE: Ontario COUNTRY: Canada ZIP: M5G 1R7 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM FC COMPATIBLE COMPUTER: PLOPS/MS-DOS SOFTWARE: Patentin Release #1.0, v
; Sequence 99, Application US/08257528B
; Patent No. 5639854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INPORMATION:
NAME: STEWART, MICHAEL I
REGISTATION UNDRER: 24,973
REFRENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEPAK: (416) 595-1163
INFORMATION FOR SEQ ID NO: 99:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 15, Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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Gaps

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US-08-465-217A-99

US-08-465-217A-99

Sequence 99, Application US/08465217A

Patent No. 5800822

GENERAL INFORMATION:

APPLICANT: SIA, Charles D.Y.

APPLICANT: CHONG, Pele

TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides

NUMBER OF SEQUENCES: 101

CORRESPONDENCE ADDRESS:

ADDRESSEE: Sim & McBurney

STRET: Suite 701, 330 University Avenue

CITY: Toronco

CONTTY: Canada

ZIP: MRG IR7

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: PERDABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: PAD-100 MGsk

CLASSIFICATION NUMBER: 08-5-17A

PRING DATE: 09-JUN-1994

PRING DATE: 09-JUN-1993

CLASSIFICATION NUMBER: 08-0-JUN-1993

CLASSIFICATION NUMBER: 08-JUN-1993

                                                                                                                                                                                                                                                                                            100.0%; Score 77; DB 1; Length 24; 100.0%; Pred. No. 1.3e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 24;
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ilarity 100.0%; Pred. No. 1.3e-05;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-486 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
  1038-487 MIS:jb
                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                       TELECOMMUNICATION INFORMATION TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                 1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                TELEFAX: (416) 595-1163
INPORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 24 amino acida
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                 15; Conservative
                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 15; Conserva
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Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: siz
                                                                                                                                                                                                                          linear
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TOPOLOGY:
US-08-463-966A-99
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US-08-465-217A-99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 77; DB 1; Length 24; 100.0%; Pred. No. 1.3e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
Patent No. 5795955
GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDERES:
ADDRESSEE: Sim & Meburney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

CLASSIFICATION NUMBER: 08/257,528

FILING DATE: 09-JUN-1994

CLASSIFICATION: 424

PRIOR APPLICATION: 424

PRIOR APPLICATION NUMBER: 08/073,378

RILING DATE: 09-JUN-1993

CLASSIFICATION: 424

RILING DATE: 09-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,602A
FILING DATE: 02-JUN-1995
CLASSIFICATION: 424
                                                                CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/257,528
FILING DATE: 09-UNA-1994
CLASSIFICATION A44
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-45
FELECOMMUNICATION: 416,595-1155
TELECOMMUNICATION: 416,595-1163
FILEPRAY: (416) 595-1163
FILEPRAY: (416) 595-1163
FILEPRAY: (416) 595-1163
FILERAY: (416) 595-1163
FILERAY: 24 AMINO ACIÐS
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NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7 RIQRGPGRAFVTIGK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 24 amino acida
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-460-602A-99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 76
US-08-463-966A-99
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100.0%; Score 77; DB 2; 100.0%; Pred. No. 1.3e-05
                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Kaiser, Annemarie
REGISTRATION NUMBER: 37,649
REFERENCE/DOCKET NUMBER: METRICS.12CPC1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-235-8550
TELEFAX: 619-235-0176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 619-215-01.0
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       internal
Horal, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 15; Conserva
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                                                                                                                                          US-08-464-329A-99
; Sequence 99, Application US/08464329A
; Pattent No. 5817754
; GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
; APPLICANT: KLEIN, Michel H.
; TTLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
; CORRESPENDENCE ADDRESS:
; ADDRESSE: Sim & McBurney
; STREET: Sunte 701, 330 University Avenue
; CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER KEALABLE FORM:
MEDIUM TYPE: FLORDY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM:
PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,329A
FILING DATE: 05-JUN 1995
CLASSIFICATION TABER: US/08/464,329A
FILING DATE: 09-JUN-1994
CLASSIFICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR BATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY/AGBNT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 24,973
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1153
INFORMATION FOR END ID NO: 99:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7 RIQRGPGRAFVTIGK 21
                       RIORGPGRAFVTIGK 15
                                                             7 RIORGPGRAFVTIGK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ESS: single
linear
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                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
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                                                                                                                               RESULT 78
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TITLE OF INVENTION: PERFIDES FOR USE IN VACCINATION AND
TITLE OF INVENTION: INDUCTION OF NEUTRALIZING ANTIBODIES AGAINST HUMAN
TITLE OF INVENTION: INMUNODEFICIENCY VIRUS
NUMBER OF SEQUENCES: 41
CORRESPONDENCE ADDRESS:
ADDRESSEE: KNOBBE, MARTENS, OLSON AND BEAR
STREET: 620 NEWPORT CENTER DRIVE 16TH FLOOR
CITY: NEWPORT BEACH
STATE: CA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KESULT 80

US-08-462-507A-99

Sequence 99, Application US/08462507A

Patent No. 5876731

GENERAL INFORMATION:

APPLICANT: CHONG, Pele

APPLICANT: KLEIN, Michel H.

TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides

NUMBER OF SEQUENCES: 101

CORRESPONDENCE ADDRESS:

ADDRESSEE: Sim & McBurney

STARE: Outerio OCOUNTRY: Canada

ZIP. ...

ZIP. ...
                                                                                                                                                                                                                                         ZIP: 92660
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: BATENIN PC-DOS/MS-DOS
SOFTWARE: PATENIN PREBABBE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/493,235
FILING DATE: 20,1010)1995
CLASSIPICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/462,507A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.36-05;
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NGS-08-73-425A-160

Sequence 160, Application US/08723425A

Sequence 160, Application US/08723425A

Sequence 160, Application US/08723425A

Sequence 160, Application

Sequence 160, Application

Sequence 160, Application

GENERAL INFORMATION:

PROCESS FOR THE DETERMINATION OF

TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT

TITLE OF INVENTION: BPITOPES AND THERE USE IN A PROCESS FOR DETERMINATION OF

TITLE OF INVENTION: ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING ...

NUMBER OF SEQUENCES: 453

CORRESPONDENCE ADDRESS:

ADDRESSEE: NIXON & VANDERHYE, P.C.

STREET: 1100 NORTH GLEBE ROAD, BTH FLOOR

CITY: Arlington

STATE: VA

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 1.3e-05; tive 0; Mismatches 0; Indels
Sequence 99, Application US/08467881A
Patent No. 5951986
GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: KIEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATIONS STSTEM:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,881A
FILING DATE: 06-JUN-1995
CLASSIPICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1994
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION NUMBER: 109-307
REFERENCE/DOCKET NUMBER: 1038-488 MIS:jb
TELECOMMUNICATION NUMBER: 1038-488 MIS:jb
                                                                                                                                                                                                                                                                                                    CITY: Toronto
STATE: Ontario
CONTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 99:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 RIQRGPGRAFVTIGK 21
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Best Local Similarity 100.0
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-467-881A-99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BICHINITATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES, PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROCESS FOR THE DETERMINATION OF PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CERDEACA.

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEFFILDE
TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTY
TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIER
TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIER
TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSIT
NUMBER OF SEQUENCES: 453
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACIBLE
COMPUTER: PREPARING SYSTEM:
COMPUTER: PREPARING SYSTEM:
COMPACIE: PATENTIN Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION NUMBER: US/08/146,028
INPORMATION FOR SEQ ID NO: 160:
SEQUENCE CHARACTERISTICS:
CALL OF THE PROCESS FOR THE DETERMINATION FOR SEQ ID NO: 160:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 1.3e-05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 77; DB 2; L
Best Local Similarity 100.0%; Pred. No. 1.3e-05;
Matches 15; Conservative 0; Mismatches 0;
                                                                                                                                                CLASSIFICATION: 444

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-UNA-1993
CLASSIPICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERRACE/DOCKET NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPROME: (416) 595-1155
TELEPRATION FOR SED ID NO: 99:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 160, Application US/08146028; Patent No. 5891640; GENERAL INFORMATION:
  FILING DATE: 05-JUN-1995
CLASSIFFCATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7 RIQRGPGRAFVTIGK 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 single
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; MOLECULE TYPE: peptide
US-08-146-028-160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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US-08-467-881A-99
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Gaps

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Sequence 14, Application US/09790497A
Patent No. 6649735
GENERAL INFORMATION:
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Patent No. 6210903
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                             1 RIQRGPGRAFVTIGK 15
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
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                                                                                            STRANDEDNESS: unknown TOPOLOGY: unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                   MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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US-09-112-206-160
                                                                                                                                                                                                                                                     ; NAME/KEY:
; LOCATION:
US-08-480-332-2
TELEFAX:
                                                                                                                                                                                                                                     FEATURE:
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Patent No. 6180134

GENERAL INFORMATION:
APPLICANT: Zalipsky, Samuel; Woodle, Martin; Martin, Francis;
APPLICANT: Barenholz, Yechezkel
TITLE OF INVENTION: Method
TITLE OF INVENTION: Method
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Absociates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/723,425A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
100.0%; Score 77; DB 3; I
Best Local Similarity 100.0%; Pred. No. 1.3e-05;
Matches 15; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Avenue, Suite 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 38,563
REFERENCE/DOCKET NUMBER: 5325-0115.31
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PEDLICATION NUMBER: US/08/480,332
RILING DATE: 7-JUN-1995
CLASSIPRCATION: 424
RRICA APPLICATION DATA:
APPLICATION NUMBER: US/08/316,436
FILING DATE: 29-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/035,443
RILING DATE: 23-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAMME: MOAL: JUNG M.
REGISTRATION NUMBER: 38,563
                                                                                                                                                                                                                                                              1487-13
                                                                                                                                                         FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
REGISTRATION NUMBER: 36,663
REFRENCE/POCKET NUMBER: 1487-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-816-4000
INFORMATION FOR SEQ ID NO: 160:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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US-08-480-332-2
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PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES, PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
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                                                                                                                                                                                                                                            Length 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 77; DB 3; I
100.0%; Pred. No. 1.3e-05;
                                                                                                                                                                                                                                               100.0%; Score 77; DB 3; 3 100.0%; Pred. No. 1.3e-05;
                                                                                                                                                                                                                                                                                                    0; Mismatches
HYPOTHETICAL: NO ANTI-SENSE: NO ORIGINAL SOURCE: INDIVIDUAL ISOLATE: Peptide 2, Fig. 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: PROCESS FOR THE TITLE OF INVENTION: CORRESPONDING TO TITLE OF INVENTION: IN A PROCESS FOR TITLE OF INVENTION: PROCESS FOR PREFITURE OF INVENTION: PROCESS FOR PREFITURE OF INVENTION: PROCESS FOR PREFITURE OF SEQUENCES: 453
COMPUTER READABLE FORM: MEDIUM TYPE: FLORPY disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICATION NUMBER: US 08/146,028
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TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
TITLE OF INVENTION: A PROCESS FOR DETERMINATION OF ANTIBODIES OF
TITLE OF INVENTION: BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
TITLE OF INVENTION: BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
TITLE OF INVENTION: CONTAINING THEM
FILE REPRENCE: 2752-11
CURRENT PILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 08/723,425
PRIOR APPLICATION NUMBER: 08/723,425
PRIOR APPLICATION NUMBER: 08/723,425
PRIOR PILING DATE: 1993-11-22
PRIOR PILING DATE: 1993-11-22
PRIOR FILING DATE: 1993-09-08
PRIOR FILING DATE: 1993-09-08
PRIOR FILING DATE: 1993-09-08
PRIOR FILING DATE: 1993-09-08
PRIOR PILING DATE: 1993-03-08
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COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/680,497
FILING DATE: 06-OCT-2000
RIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/146,028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 77; DB 4; I 100.0%; Pred. No. 1.3e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        , ORGANISM: Human immunodeficiency virus US-09-576-824A-160
                                                                                                                                                                                                                                                                                                     Sequence 160, Application US/09576824A Patent No. 6667387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 160, Application US/09680497; Patent No. 6709828; GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 160:
SEQUENCE CHARACTERISTICS:
                 1 RIQRGPGRAFVTIGK 15
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TITLE OF INVENTION: PROC
TITLE OF INVENTION: IN A
TITLE OF INVENTION: PEPT
TITLE OF INVENTION: PEPT
TITLE OF INVENTION: PROC
NUMBER OF SEQUENCES: 453
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                 8 RIORGPGRAFVTICK
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                      US-09-576-824A-160
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Facture 160, Application US/09790497A

Facture 160, Application US/09790497A

Facture 160, Application US/09790497A

FACTURE OF INVENTION:

FITLE OF INVENT
        PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR USE IN
A PROCESS FOR DETERMINATION OF ANTIBODIES OF
BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
EPITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
CONTAINING THEM
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TITLE OF INVENTION: PROCESS FOR THE DETERMINANT TITLE OF INVENTION: A PROCESS FOR DETERMINANT TITLE OF INVENTION: BIOTINYLATED PETERMINANT TITLE OF INVENTION: BIOTINYLATED PETERMINANT TITLE OF INVENTION: EPITOPES, A PROCESS FOR FITTLE OF INVENTION: EPITOPES, A PROCESS FOR FITTLE OF INVENTION: EPITOPES, A PROCESS FOR FORTEN TO FILL OF INVENTION: EPITOPES, A PROCESS FOR FITTLE REFERENCE: 2752-16

CURRENT APPLICATION NUMBER: 09/756,824

PRIOR FILING DATE: 2000-05-23

PRIOR FILING DATE: 1996-09-30

PRIOR FILING DATE: 1993-11-22

PRIOR FILING DATE: 1993-11-22

PRIOR FILING DATE: 1993-10-08

PRIOR FILING DATE: 1993-00-88

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Best Local Similarity 100.0
Matches 15; Conservative
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US-09-790-497A-160
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512-474-7577
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 10301 SCITY: Houston STATE: Texas COUNTRY: USA ZIP: 77025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                            Texas
                                                                                                                                                                                                                                                                     RY: US
77210
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                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                         STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 92
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                                                                                                                                                               Gaps
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GENERAL INFORMATION:
APPLICANT: THE ROCKEFELLER UNIVERSITY
TITLE OF INVENTION: MULTIPLE ANTIGEN PEPTIDES FOR USE AS HIV
TITLE OF INVENTION: WACTINES
NUMBER OF SEQUENCES: 21
CORRESPONDENCES. 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0%; Pred. No. 1.3e-05;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                         Length 24;
                                                                                                      Query Match 100.0%; Score 77; DB 4; Length 24 Best Local Similarity 100.0%; Pred. No. 1.3e-05; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTER: Massacinsetts

ZIP: 02110-2804

COMPUTER READABLE FORM:
MEDIUNT TYPE: 3.5" Diskette, 1.44 Mb

COMPUTER: IBM PS/2 Model 502 or 558X

ODERATING SYSTEM: IBM P.C. DOS (Version 3.30)

SOFTWARE: WordPerfect (Version 5.1)

CURRENT APPLICATION NUMBER: PCT/US92/06688

FILING DATE: 19920011

CLASSIFICATION: 424

FRICA REPLICATION NUMBER: 744,281

FILING DATE: 13 August 1992

ATTORNEY/ASETY INVERMATION:
NAME: Paul T. Clark

REFERENCE/DOCKET NUMBER: 30,162

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GENERAL INFORMATION:
APPLICANT: BOARD OF REGENTS, THE UNIVERSITY OF
APPLICANT: TEXAS SYSTEM
APPLICANT: SASTRY, Jagannadha K.
APPLICANT: ARLINGHAUS, Ralph B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                     8 RIORGPGRAFVTIGK 22
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
; TOPOLOGY: linear; MOLECULE TYPE: peptide US-09-680-497-160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Massachusetts
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Sequence 1, Application US/07950571A
PATENT NO. 585400
GENERAL INFORMATION:
APPLICANT: Chang, Tee Wen, Fung, Michael S.C., Sun, Bill N.C., Sun, Cecily R.Y.,
APPLICANT: Chang, Nancy T.
TITLE OF INVENTION: Monoclonal Antibodies which Neutralize HIV-1 Infection
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSES: Tanox Biosystems, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
APPLICANT: PLATSOUCAS, Chris D.
APPLICANT: NEHETE, Pramod N.
ITILE OF INVENTION: METHODS AND COMPOSITIONS
TITLE OF INVENTION: FOR ELICITING IMMUNE OR ANTI-INFECTIVE RESPONSES
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
STREET: P.O. Box 4433
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 77; DB 5; I
100.0%; Pred. No. 1.3e-05;
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MEDIUM TYPE: 3.5" Hi Denaity Diskette
COMPUTER: IBM PS/2
OPERATING SYSTEM: DOS, Version 3.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Parker, David L. REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: UTFC305PCT
TELECOMMINICATION INFORMATION:
TELEPHONE: 512-320-7200
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10378
FILING DATE: 19921202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 07/800,932
FILING DATE: December 2, 1991
PRIOR APPLICATION NUMBER: 07/945865
FILING DATE: September 16, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Tanox Biosystems, Inc. 10301 Stella Link Rd.
                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMEDIUM: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: Not Applicable INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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Gaps

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Length 25;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: BATENIE: PC-DOS/MS-DOS
SOFTWARE: PATENIE: PC-DOS/MS-DOS
SOFTWARE: PATENIE: PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,324
FILING DATE: 07-JUN-1995
PRIOR APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAMME: BVARA: BATTY
REGISTRATION NUMBER: 370132-2000
TELECOMMUNICATION NUMBER: 370132-2000
TELECOMMUNICATION NUMBER: 370132-2000
                                                                                                                                                                                                                                                                                         100.0%; Score 77; DB 2; 1 100.0%; Pred. No. 1.4e-05;
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                                                                                                                                                                                                                                                                                                                                    Mismatches
                     03495.0088-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Curtis, Morris, & Safford ADDRESSEE: C/O Barry Evans STREET: 530 Fifth Avenue CITY: New York STATE: New York COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13, Application US/08485324
Patent No. 6043093
GENERAL INFORMATION:
APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
REGISTRATION NUMBER: 25,146
REPRENCE/DOCKET NUMBER: 0349
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4132
INFORMATION FOR SEQ 10 NO: 1:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query.Match 100.0
Best Local Similarity 100.0
Matches 15, Conservative
                                                                                                                                             LENGTH: 25 amino acida
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                            not relevant
                                                                                                                                                                                                                                                                                                            Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                            TOPOLOGY: not relevant MOLECULE TYPE: peptide US-08-266-448-1
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                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                         Query Match
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INDUCTION OF PROTECTION AGAINST VIRAL
INFECTION BY SYNERGY BETWEEN VIRUS ENVELOPE GLYCOPROTEIN
AND PEPTIDES CORRESPONDING TO NEUTRALIZATION EPITOPES OF
THE GLYCOPROTEIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYRE: PLOPY disk
COMPUTER READABLE FORM:
MEDIUM TYRE: Flopy disk
COMPUTER: IBM PC compatible
COMPUTER: BM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/266,448
FILING DATE: 28-UN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/145,664
FILING DATE: 28-OCT-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/782,241
FILING DATE: 28-OCT-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/672,647
FILING DATE: 19-MAR-1991
FILING DATE: 19-MAR-1990
ATTORNEY AGENT INFORMATION:
NAME: MEYERS, KENDETH J.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 77; DB 2; Ler
Pred. No. 1.4e-05;
                                                     AFFLICATION NUMBER: US/U/950,5/IA
FILING DATE: 1992092
CLASSIFICATION: 435
PRIOR AFPLICATION: 435
PRIOR AFFLICATION DATA: AFFLICATION NUMBER: NO. 5854400 07/767,533
FILING DATE: 09/26/1991
ATTORNEY/AGENT INFORMATION:
NAME: MICABEL, Eric P.
REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: 31,211
FELECOMMUNICATION INFORMATION:
TELEPHONE: 713-664-2288
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
                                             US/07/950,571A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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Best Local Similarity 100.0%;
Matches 15; Conservative 0
  SOFTWARE: Wordperfect 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/07/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11 RIORGPGRAFVTIGK 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                             AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                    Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: D.C.
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                     US-07-950-571A-1
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Length 25;
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                 COMPUTER: FLODEY disk
MEDIUM TYPE: Floppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: Plan PC Compatible
COMPUTER: Plan PC COMPUTER:
COMPUTER: Plan PC COMPUTER:
COMPUTER: Plan PC COMPUTER:
COMPUTER: Plan PC COMPUTER:
APPLICATION NUMBER: US 08/447,506
FILING DATE: 23-MAY-1995
CLASSIFICATION NUMBER: US 08/235,437
FILING DATE: 13-MAR-1994
APPLICATION NUMBER: US 08/852,412
FILING DATE: 16-MAR-1992
ATOMNEY AGENT INFORMATION:
NAME: Evans Barry
REGISTRATION NUMBER: 22,802
REGISTRATION NUMBER: 22,802
REGISTRATION NUMBER: 22,802
REGISTRATION NUMBER: 22,802
REFERENCE/POCKET NUMBER: 370132-2000
TELEPPAN: (212) 840-3333
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LEMETHER COMPUTER: 25-MAIN ACIDS
LEMETHER CHARACTERISTICS:
LEMETHER CHARACTERISTICS:
LEMETHER COMPUTER: 25-MAIN ACIDS
LEMETHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: Ploppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COPEWARIG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/447,506
FILING DATE: 23-MAY-1995
CLASSIFICATION 1435
PRIOR APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
100.0%; Score 77; DB 3; I
Best Local Similarity 100.0%; Pred. No. 1.4e-05;
Matches 15; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVERTION:
SELECTION METHODS
INFORMATION:
ADDRESSEE: Curits, Morris, & Safford
ADDRESSEE: C/O Barry Evans
STREET: 530 Fifth Avenue
CITY: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Evans, Barry
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 31, Application US/08447506
Patent No. 6066499
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10036
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-08-447-506-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-447-506-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Elb PC compatible
COMPUTER: Elb PC compatible
COMPUTER: Elb PC compatible
COMPUTER: Datentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,324
FILING DATE: 07-JUN-1995
RICH APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: EVANB, BAITY
REGISTRATION NUMBER: 22 807
REGISTRATION NUMBER: 22 807
REGISTRATION NUMBER: 23 807
REGISTRATION NUMBER: 22 807
REGISTRATION NUMBER: 23 807
                                                                                                                                                                                   Sequence 31, Application US/08485324
Sequence 31, Application US/08485324
Sequence 31, Application US/08485324
Sequence 31, Mohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: C/O Barry Evans
STREET: S30 Fifth Avenue
CITY: New York
STATE: New York
STATE: New York
COUTTY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Curtis, Morris, & Safford
ADDRESSEE: c/o Barry Evans
STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 370132-2000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13, Application US/08447506
Patent No. 6066499
GENERAL INFORMATION:
APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 RIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 25 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SS: single
linear
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STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10036
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US-08-447-506-13
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US-08-235-437-31
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                                                                                                                                                                                                                                                              Query Match
100.0%; Score 77; DB 3; Length 25;
Best Local Similarity 100.0%; Pred. No. 1.4e-05;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 77; DB 3; Length 25; 100.0%; Pred. No. 1.4e-05; tive 0; Mismatches 0; Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-US-13-13

Sequence 13, Application US/08215437

Patent No. 6087177

GENERAL INFORMATION:
APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS

NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: CALTIS, Morris, & Safford
ADDRESSEE: CALTIS, Morris, & STATE: New York
COUNTRY: USA
ZIP: 10036
COMPUTER: EBADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/235,437
FILING ANTE: 29-APR-1994
CLASSIFICATION: ADDRESSEE
COUNTRY: APPLICATION NUMBER: US/08/235,437
FILING ANTE: 29-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22,802
IER: 370132-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
; LELEPAX: (212) 840-333

FELEPAX: (212) 840-333

FINFORMATION FOR SEQ ID NO: 31:

SEQUENCE CHARACTERISTICS:

LENGTH: 25 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Evans, Barry
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                          RIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 25 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                      US-08-447-506-31
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RESULT 99

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Gaps
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                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: Floppy disk

COMPUTER: Elby PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/235,437

FILING DATE: 29-APR-1994

CLASSIFICATION: 435

FILING DATE: 19-APR-1994

APPLICATION NUMBER: US 07/852,412

FILING DATE: 16-MAR-1992

ATTORNEY/AGENT INFORMATION:

NAME: EVAND: BALTY

REGISTRATION NUMBER: 22,802

REFERENCE/DOCKET NUMBER: 370132-2000

TELEPHONE: (212) 840-3133

TELEPHONE: (212) 840-0712

INFORMATION POR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:

LENGTH: 25 amino acids

TVODE: LENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 77; DB 3; I
Sequence 31 Application US/08235437
Patent No. 6087177
GENERAL INFORMATION:
APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: C/O Barry Evans
STREET: S30 Pifth Avenue
CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 13, Application US/08447515

Patent No. 6162640

GENERAL INFORMATION:
TITLE OF INVENTION: SELECTION METHODS

NUMBER OF SEQUENCES: 31

CORRESPONDENCE ADDRESS:
ADDRESSEE: C/O Barry Evans
STREET: 530 Fifth Avenue
CITY: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10036
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100.0%; Score 77; DB 4; I
100.0%; Pred. No. 1.4e-05;
                                                                      100.0%; Score 77; DB 3; I
100.0%; Pred. No. 1.4e-05;
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                                                                                                         Mismatches
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Patent No. 5932218
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                                                                                                                                                1 RIORGPGRAFVTIGK 15
                                                                                                                                                                         8 RIQRGPGRAFVTIGK 22
                                                                        Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 15; Conservative
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
CRGANISM: Homo sapiens
US-09-593-870A-31
               ; TOPOLOGY: linear
US-08-447-515-31
                                                                                                                                                                                                                                         RESULT 102
US-09-593-870A-31
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                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 77; DB 3; Length 25; 100.0%; Pred. No. 1.4e-05; tive 0; Mismatches 0; Indels
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MEDIUM TYPE: RADABLE FORM:

MEDIUM TYPE: RADABLE FORM:

COMPUTER: IBM PC compatible

SOFTWARE: Patentin Release #1.0, Version #1.25

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/447,515

FILING DATE: 23-MAY-1995

APPLICATION NUMBER: US 08/235,437

FILING DATE: 16-MAR-1994

APPLICATION NUMBER: US 07/852,412

FILING DATE: 16-MAR-1992

ATTORNEY/AGENT INFORMATION:

NAME: EVANS BALTY

REGISTRATION NUMBER: 22,802

REGISTRATION NUMBER: 23,370132-2000

TELECHOME: (212) 840-3333
           PILLING DATE: 23-MAY-1995
CLASSIFICATION: 435
PRIOR APPLICATION 1945
APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Evans, Barry
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
TELEPHONE: (212) 840-3333
TELEPHONE: (212) 840-3333
TELEPHONE: (212) 840-313;
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids

"""""" and acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/447,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8 RIQRGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                                                                                                                                                                                                                              single
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                            , TOPOLOGY: linear
US-08-447-515-13
                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
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APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Berzofsky, Jay A.
APPLICANT: Berzofsky, Jay A.
APPLICANT: Barzof C. D.
APPLICANT: Brian Mushuncri
APPLICANT: Shirah Mushuncri
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE BITCH, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STRATE: Virginia
COUNTRY: USA
CITY: Falls Church
STRATE: Poppy disk
COMPUTER ERADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATION SYSTEMS: PetentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
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                                                      Gaps
                                                                                                                                                                                                                                                                                          Sequence 31, Application US/09593870A

| Sequence 31, Application US/09593870A
| Patent No. 6548643
| GENERAL INFORMATION:
| APPLICANT: MCKENZIE, Ian F.C.
| APPLICANT: Postcologoulos, Vasso
| APPLICANT: Pteteraz, Geoff Allan
| TITLE OF INVENTION: Use in Immunotherapy
| FILE REPRENCE: 2368-MCKenzie
| CURRENT APPLICATION NUMBER: US/09/593,870A
| CURRENT FILING DATE: 1090-06-14
| PRIOR FILING DATE: 1998-12-30
| NUMBER OF SEQ ID NOS: 69
| SOFTWARE: FastSEQ for Windows Version 3.0
| SEQ ID NO 31
| LENGTH: 25
| CURRENT PROBLEM NUMBER: US/09/223,043
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 25;
Length 25;
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RESULT 105

US-060-988A-12

US-060-988A-12

Sequence 12, Application US/08060988A

Sequence 12, Application US/08060988A

Sequence 12, Application US/08060988A

Sequence 12, Application US/08060988A

APPLICANT: Berzofsky, Jay A.

APPLICANT: Pendleton, C. David

APPLICANT: Nara, Peter

APPLICANT: Shirai, Mutsunori

TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES

TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND

TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND

TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1

NUMBER OF SEQUENCES: 48

CORRESPONDENCE ADDRESS:

ADDRESSEE: Fish & Richardson P.C.

STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 96.1%; Score 74; DB 3; Length 15; Best Local Similarity 93.3%; Pred. No. 2.5e-05; Matches 14; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRETTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: PassESEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION NUMBER: 07/847,311
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY-AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: 08830/022001
TELEBHONE: 617/542-5070
                    NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022003
TELECOMMUICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 200154
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RIQKGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                              TELEX: 2017/11 050
TELEX: 20154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-455-685-12
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APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Shiral, Muteunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
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Pred. No. 2.5e-05;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: Peptide
LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-4, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTEY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDLIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: BEASTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/45,685
FILING DATE: 31-WAY-1995
APPLICATION NUMBER: 08/060,988
FILING DATE: 14-WAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-WAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY ABGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 1173-434P
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEFAX: 703-205-8000
TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: Amino acids
                                                                                                                                                                                                                                     1173-434P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Fish & Richardson P.C.
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      . Sequence 12, Application US/08455685; Parent No. 6214347; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 93.3%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIQKGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
FRACMENT TYPE: internal
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 225 F
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
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STREET:
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ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
LYMPHOCYTES AGAINST HIV
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                                                                                            Query Match 96.1%; Score 74; DB 3; Length 15; Best Local Similarity 93.3%; Pred. No. 2.5e-05; Matches 14; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 22040-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: 1.75
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "pl8-4, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
STADRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
CITY: Falls Church
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRICR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MA-1993
ATTORNEY AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-20-8000
                                                                                                                                                                                                                                                                                                                     Sequence 12, Application PC/TUS9405142
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: ELECTING NEUTR
TITLE OF INVENTION: LYMPHOCYTES AGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 12: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                           1 RIOKGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide FRAGMENT TYPE: internal
               TOPOLOGY: linear

MOLECULE TYPE: peptide

US-08-060-988A-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
amino acid
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Matches 14; Conserv
                                                                                                                                                                                                                                                                                       RESULT 106
PCT-US94-05142-12
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RESULT 107

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                                               APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. D.
APPLICANT: Pendleton, C. D.
APPLICANT: Para, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Patent No. 593218
GENERAL INFORMATION:
APPLICANT: Berior Seffrey D.
APPLICANT: Pendleton, C. D.
APPLICANT: Nara, Peter
APPLICANT: Shirai, Matsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label= peptide
/note= "p18-9, see Table V"
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93.3%; Pred. No. 3.6e-05;
cive 0; Mismatches 1
                                                                                                                                                                                                                                                                                ADDRESSEE: Birch, Stewart, Kolasch & Birch STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAX-1993
ATTORNEY/AGENT INFORMATION:
Sequence 17, Application US/08455625
Patent No. 593218
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE DOCKET NUMBER: 1:
TELECOMMUNICATION INFORMATION
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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CTHER INFORMATION:
CTHER INFORMATION:
US-08-455-625-17
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Matches 14; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 22040-0747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                  CITY: Fal
STATE: Vi
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 108
US-08-455-625-23
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SSE: Fish & Richardson P.C. 225 Franklin Street Boston
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1 Similarity 93.3%;
14; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 15 amino acids
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ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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Best Local Similarity
Matches 14; Conserv
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CITY: Bo
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TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS: 36
CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
COUNTRY: USA
ZIP: 22040-0747
COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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US-08-455-685-17

Sequence 17, Application US/08455685

Patent No. 6214347

GENERAL INFORMATION:
APPLICANT: Berzefsky, Jay A.
APPLICANT: Alersy Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Shirai, Mutsunori
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: WILPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: Peptide
LOCATION: 1.15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-15, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94.8%; Score 73; DB 2; 93.3%; Pred. No. 3.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA.
PRIOR APPLICATION UDBER: US 08/060,988
FILING DATE: 14-MAY.1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ 10 NO: 23:
SUGURNE TO 3-205-8050
INFORMATION FOR SEQ 1D NO: 23:
SUGURNE TARREFERENCE TO 3-205-8050
INFORMATION FOR SEQ 1D NO: 23:
SUGURNE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              :B: Fish & Richardson P.C.
225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 93.3<sup>1</sup>
...han 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-455-625-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
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Sequence 23, Application US/08455685

Sequence 23, Application US/08455685

Patent No. 6114347

GENERAL INFORMATION:
APPLICANT: Berzofeky, Jay A.
APPLICANT: Berzofeky, Jay A.
APPLICANT: Anlers, Jeffrey D.
APPLICANT: Nara, Peter
APPLICANT: Shiral, Mutenori
APPLICANT: Shiral, Mutenori
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 73; DB 3; Length 15;
Pred. No. 3.6e-05;
0; Mismatches 1; Indels
COMPUTER: Diskette
COMPUTER: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEG for Windows Version 2.0
SURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
APPLICATION NUMBER: 07/847,311
FILING DATE: 14-MAY-1992
APPLICATION NUMBER: 07/847,311
FILING DATE: 25-AUG-1991
APPLICATION NUMBER: 07/146,692
FILING DATE: 26-AUG-1991
APPLICATION NUMBER: 07/146,692
FILING DATE: 26-AN-1988
ATTORNEY/AGENT INFORMATION:
NAME: BEATLE, INGINA A.
REFERENCE/DOCKET NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: B-42,306
TELECHOME: 617/642-5070
TELECHOME: 617/642-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
PRILING DATE: 31-MAY-1995
PRIOR APPLICATION DATA:
```

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Sequence 23, Application US/08060988A Patent No. 6294322 GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A. APPLICANT: Ahlers, Jeffrey D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Berzofeky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                            TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 93.3%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 RIORGPGRVFVTIGK 15
                                                                                                                                                                                                                                   , TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-060-988A-17
                                                                                                                              TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                 617/542-50'
617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-060-988A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
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                                                             TELEPHONE:
TELEFAX: 61
TELEX: 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -08-060-988A-23
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APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AN TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94.8%; Score 73; DB 3;
93.3%; Pred. No. 3.6e-0
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REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022001
                                                                                                                                                                                                                                                                                                                                                                                    08830/022003
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-ANN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/0
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: 200154
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LEBNGTH: 15 amino acide
TELENGTH: 15 amino acide
TELENGTH: 15 amino acide
TELENGTH: 15 amino acide
TELENGTH: 15 amino acide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 17, Application US/08060988A ; Patent No. 6294322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 94.8
Best Local Similarity 93.3
Matches 14; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , MOLECULE TYPE: peptide US-08-455-685-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Ri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION
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TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AN
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
CARRESPONDENCE ADDRESS:
ADDRESSEE: Pish & Richardson P.C.
STREET: 225 Franklin Street
                                                            ö
  Length 15;
                                                         1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: MA
COMPUTER: MA
COMPUTER: TADABLE FORM:
MEDLUM TYPE: Diskette
COMPUTER: IBM Compartible
OPERATING SYSTEM: Windows95
SOFFWARE: FastSEQ for Windows Version 2.0
SURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 05-MAR-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY-AGENT INFORMATION:
NAME: BEALTISH INFORMATION:
NAME: BRATSTRATION NUMBER: P+42.306
Score 73; DB 3;
Pred. No. 3.6e-05;
0; Mismatches 1
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Query Match
Best Local Similarity 93.3
Matches 14; Conservative
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MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: Peptide
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                                                                                                                                                                                                                       Sequence 17. Application PC/TUS9405142

GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STRET PO. Box 747
CITY: Falls Church
STATE: Virginia
COUNTY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                               Gape
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94.8%; Score 73; DB 5; Length 15;
Best Local Similarity 93.3%; Pred. No. 3.6e-05;
Matches 14; Conservative 0; Mismatches 1; Indels
  Length 15;
                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: 1.715
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-9, see Table V"
  Score 73; DB 3; Pred. No. 3.6e-05;
                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
APLICATION NUMBER: US 08/060,988
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
RECISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8050
TELEPHONE: 703-205-8050
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCT-US94-05142-23; Sequence 23, Application PC/TUS9405142; GENERAL INFORMATION:
Query Match
Best Local Similarity 93.3%;
Matches 14; Conservative 1
                                                                                        1 RIQRGPGRAFVTIGK 15
                                                                                                                                 1 RIORGPGRAFVTIGO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIQRGPGRVFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : 15 amino acids
amino acid
GY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCT-US94-05142-17
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В
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TITLE OF INVESTION: CHOCKETTS SYMPHETE REPTIDE CONSTRUCT
TITLE OF INVESTION: INVESTION: INVESTION ELICITING MENTALIZING ANTIBODIES AND CTROTOKIC I TITLE OF INVESTION: INVESTIGATION: INVESTION: INVESTIGATION: INVEST
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Sequence 9, Application US/08060988A
; Sequence 9, Application US/08060988A
; Patent No. 6294322
; GENERAL INFORMATION:
; APPLICANT: Berescieky, Jay A.
; APPLICANT: Anlers, Jeffrey D.
; APPLICANT: Anlers, Jeffrey D.
; APPLICANT: Anlers, Deference
; APPLICANT: Anlers, MULTIDETERMINANT PEPTIDES
; TITLE OF INVENTION: THAT ELICIT
; TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE
; TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE
; TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE
; TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
; TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
; TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
; TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE
; CONTRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson P.C.
; STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93.5%; Score 72; DB 3; Length 14; 100.0%; Pred. No. 4.8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: EBM Compatible
COMPUTER: EBM Compatible
OPERATING SYSTEM: Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
FILING DATE: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-4UG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1981
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: HARTHER TANGER TOWNEY AND THE TOWNEY AND TOWNEY AND THE 
                            FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: BEALIE, Ingrid A.
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: 08830/022003
TELEPHONE: 617/542-8906
TELEPHONE: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Prec. ...
08/090,988
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REGISTRATION NUMBER: P-42,306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Beattie, Ingrid A.
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Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , MOLECULE TYPE: peptide US-08-455-685-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM
   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-060-988A-9
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APPLICANT: BALEZGESKY, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendateron, C. David
APPLICANT: Nata, Peter
APPLICANT: Shiral, Mucaunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                   OPERATION STRICTS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
TILING DATE:
FILING DATE:
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14 MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svenseon, Leonard R.
REGISTRATION NUMBER: 30330
REBERRENGE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: US
ZIF: 02110-2804
ZIF: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEC for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION UNDERS: US/08/455,685
FILING DATE: 31-MAY'1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
93.5*; Score 's, --
Best Local Similarity 100.0%; Pred. No. 4.8
Matches 14; Conservative 0; Mismatches
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225 Franklin Street
       ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9, Application US/08455685
Patent No. 6214347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 IQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-455-685-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 116
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SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NOMBER: PCT/US95/03236
PILING DATE: 13-MAR-1995
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 72; DB 5; I Pred. No. 4.8e-05
                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Imbra, Richard J.
REGISTRATION NUMBER: 37,643
REFERENCE/DOCKET NUMBER: FP-SI 1394
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                      ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                             2 IORGPGRAFVTIGK 15
                                                                                                                                                1 IORGPGRAFVTIGK 14
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Best Local Similarity 100.
Matches 14; Conservative
                                                                        14; Conservative
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MOLECULE TYPE: peptide
PCT-US95-03236-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
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                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: BLICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STRET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 22040-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: 1..14
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "pl8-1, see Table V"
                                                                                                                                                                                                                                                                                                                                                     4.8e-05;
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100.0%; Pred. No. 4.8e-05;
                                                                                                                                                                                                                                                                                                                                  Score 72; DB 3;
                                                                      08830/022001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISCRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9, Application PC/TUS9405142 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                93.5%; Scc.
100.0%; Pred
0; N
REFERENCE/DOCKET NUMBER: 0883v
TELEPHONE: 617/542-5070
TELEPHONE: 617/542-8006
; TELEEX: 200154
; INPORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
; LENGTH: 14 amino acids
TYPE: amino acid
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                             2 IORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 IORGPGRAFVTIGK 14
                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: 703-205-8050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                TOPOLOGY: linear
HOLECULE TYPE: peptide
US-08-060-988A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: Peptide LOCATION: 1..14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ropology: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 118
PCT-US94-05142-9
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GENERAL INFORMATION:
APPLICANT: University of Southern California
TITLE OF INVENTION: Methods to Diagnose and Treat HIV-1
TITLE OF INVENTION: Infection
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSER: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
COUNTRY: USA
                                                                                                                                                                                                              Sequence 29, Application PC/TUS9503236
GENERAL INFORMATION:
APPLICANT: University of Southern California
TITLE OF INVENTION: Methods to Diagnose and Treat HIV-1
TITLE OF INVENTION: Infection
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
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Sequence 19, Application US/08455625

Patent No. 5932218

GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A.
APPLICANT: Allers, Jeffrey D.
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYT
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
MUMBER OF SEQUENCES: 36
COURDED PROMISE AND COURT OF AUTHORITY AND CYTER
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                                                                            Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STRET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
COUNTRY: USA
ZIP: 22040-0744
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC Compatible
OPERATION SYSTEM: PC-005/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MX-1993
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 31330
RELEFRANICATION NUMBER: 31330
RELEFRANICATION NUMBER: 31330
RELEFRANICATION NUMBER: 1173-434P
TELEFRANICATION NUMBER: 1173-434P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , LUCATION: 1..15
; OTHER INFORMATION: /label= peptide
; OTHER INFORMATION: /note= "p18-11, see Table V"
US-08-455-625-19
                                                                                Score 72; DB 1; L. Pred. No. 5.2e-05;
                                                          93.5%; Sco...
100.0%; Pred. No. s...
'... 0; Mismatches
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                                                                                                                                                                                                                                                                                                2 RIQRGPGRAFVTIG 15
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Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: Peptide
              US-08-704-170-72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ð
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Sequence 72, Application US/08704170

SENERAL INFORMATION:
APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
INVENTION: MADDRESSEE: RObbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite 500
CITY: Los Angeles
STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
0
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,170
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/03236
FILING DATE: 13-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 72; DB 5; L4
Pred. No. 4.8e-05;
; Mismatches 0;
                                                                                                                                                                                                                       ATTORNEY AGENT INFORMATION:
NAME: Imbra, Richard J.
REGISTRATION NUMBER: 37,643
REFERENCE DOCKET NUMBER: FP-SI 1394
TELECOMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9001
INFORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920-331
TELECHMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 72:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 93.5%; Sc
Best Local Similarity 100.0%; P
Matches 14; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIORGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               QUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
GY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              U.S.A.
                                                                                                                                                            FILING DATE: 1
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90012
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: CA
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TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSED: Pish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 72; DB 2; Length 15;
Pred. No. 5.2e-05;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
FILING DATE: US/08/455,625
PRIOR APPLICATION NUMBER: US/08/060,988
FILING DATE: 14-MAY-1993
ATTONEY/AGENT INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-13, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Svensen, Leonard R.
REGISTRATION NUMBER: 30330
REPERRICE/DOCKET NUMBER: 1173-434P
TELEPHONE: 703-205-8000
TELEPAX: 703-205-8050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19, Application US/08455685; Sequence 19, Application US/08455685; Patent No. 6214347; GENERAL INFORMATION: APPLICANT: Beracofsky, Jay A. APPLICANT: Pahlers, Jeffrey D. APPLICANT: Pendleton, C. David APPLICANT: Nara, Peter APPLICANT: Nara, Peter APPLICANT: Shirai, Mutsunori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 93.3.
Best Local 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-455-685-19
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CITY: Bo
STATE: M
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                                             Sequence 20, Application US/08455625

Patent No. 593218

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Berzofeky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. D.
APPLICANT: Phase and Applicant: Applicant: Applicant: Composite Syntheric Construct
ITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
ITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
ITLE OF INVENTION: LIMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPOTER: IBM PC compatible
COMPOTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION BATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFRMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: 1.15

; OTHER INFORMATION: /label= peptide

; OTHER INFORMATION: /note= "p18-12, see Table V"

US-08-455-625-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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US-08-455-625-21
Sequence 21, Application US/08455625
Fatent No. 5932218
GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A.
APPLICANT: Anlers, Jeffrey D.
APPLICANT: Pendleton, C. D.
APPLICANT: Nara, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: SVENBBON, LEGNARD R.
REGISTRATION NUMBER: 30330
REPERENCE/DOCKET NUMBER: 1173
TELECOMMUNICATION INFORMATION:
TELEFAX: 703-205-8050
TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: 11 inear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIORGPGRAFVAIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 14; Conserv
                                     US-08-455-625-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ò
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Sequence 21, Application US/08455685
; Sequence 21, Application US/08455685
; Patent No. 6214347
; GENERAL INFORMATION:
    APPLICANT: Beriofsky, Jay A.
    APPLICANT: Ahlers, Jeffrey D.
    APPLICANT: Pendleton, C. David
    APPLICANT: Nara, Peter
    APPLICANT: Nara, Mutriberentines THAT ELICIT
    TITLE OF INVENTION: MUTLIBER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE
    TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE
    TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
    NUMBER OF SEQUENCES: 40
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Fish & Richardson P.C.
    STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 72; DB 3; Length 15;
Pred. No. 5.2e-05;
0; Mismatches 1; Indels
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ZIP: 02110-2804

COMPUTER READABLE FORM:
MEDUTUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/060,988
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/847,311
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
APPLICATION NUMBER: 07/148,692
               APPLICATION NUMBER: 08/060,988
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/ARENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022003
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIORGPGRAFVAIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , MOLECULE TYPE: peptide US-08-455-685-20
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY:
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 127
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| Sequence 20, Application US/08455685
| Patent No. 6214347
| SENERAL INFORMATION:
| APPLICANT: Berzofsky, Jay A. APPLICANT: Pendleton, C. David
| APPLICANT: Pendleton, C. David
| APPLICANT: APPLICANT: Mutsunori
| TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
| NUMBER OF SEQUENCES: 40
| CORRESPONDENCE ADDRESS: A ADDRESSE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
0
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Pred. No. 5.2e-05;
0; Mismatches 1; Indels
                  COUNTIEST US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette COMPUTER: Diskette COMPUTER: Enable FORM:
MEDIUM TYPE: Diskette COMPATIONE
CORRATING SYSTEM: Windows95
SOFTWARE: FASTERO for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
APPLICATION NUMBER: 08/060,988
FILING DATE: 06-WAR-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-WAR-1992
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INPORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: 0942,306
REGISTRATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INPORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: 08830/022003
TELEPHONE: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: US
ZIP: 0210-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: FastSEQ for Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Fish & Richardsc
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELERA: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIORGPGRAFYTIGK 15
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Best Local Similarity 93.3.
Best Local 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide US-08-455-685-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-455-685-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE:
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Gaps

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Joseph 100 - 988A - 20

Sequence 20, Application US/08060988A

Patent No. 6294322

Patent No. 6294322

APPLICANT: Allers, Jay A.

APPLICANT: Allers, Jeffrey D.

APPLICANT: Pendleton, C. David

APPLICANT: Pendleton, C. David

APPLICANT: Nara, Peter

APPLICANT: Shirai, Allers, Al
                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                       Length 15;
                                                                                                                                                                              1, Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: NA
COUNTRY: NA
COMPUTER: D10-2804
COMPUTER: READABLE FORM:
MEDIUM TYPE: D1skette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FESTEM: Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: US/08/060,988A
FILING DATE: 05-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                       Score 72; DB 3;
Pred. No. 5.2e-05;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08830/022001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-42,306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 0
TELECOMMUNICATION INFORMATION
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                       93.5%;
                                                                                                                                                                                                                                                                       1 RIQRGPGRAFYTIGK 15
                                                                                                                                                                                                                                            1 RIQRGPGRAFVTIGK 15
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acide
                                                                                                                    Query Match
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 93.3
Matches '14; Conservative
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MOLECULE TYPE: peptide
US-08-060-988A-20
                        MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
                                              ; MULECULE 115.
US-08-060-988A-19
            TOPOLOGY:
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APPLICANT: Belzósky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: BELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: BELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
WUMBER OF SEQUENCES: 48
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ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette COMPUTER: Diskette COMPUTER: Diskette COMPUTER: Diskette COMPUTER: Diskette COMPUTER: Diskette Windows95
SOFTWARE: PRESENCE FOR Windows Version 2.0
CURRENT APPLICATION DATA: APPLICATION NOWBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 28-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42.306
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 72; DB 3; Pred. No. 5.2e-05;
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                              P-42,306
ER: 08830/022003
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 19, Application US/08060988A; Patent No. 6294322
NAME: Beattie, Ingrid A.
REGISTATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 0883
TELEPHONE: 617/542-5070
TELEPHONE: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 21: SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acids
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INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                93.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RIORGPGRAFVTTGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 93.3
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-455-685-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
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PCT-US94-02631-72
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                                           GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: HILPETERMINANT PEPTIDES
TITLE OF INVENTION: HELPER T.LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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APPLICANT: Takehana, Yoshi
APPLICANT: Takehana, Yoshi
APPLICANT: Threamann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 72; DB 3; Length 15;
Pred. No. 5.2e-05;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTY: US
ZIP: 02110-2804

COMPUTER READABLE FORM:
MEDIUM TYPE: Disketter
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILNG DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILNG DATE: 26-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 26-ANG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-AN-1988
ATTORNEY/AGENT INFORMATION:
NAME: BEATLE, INGIA A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: B-42,306
REFERENCE/DOCKET NUMBER: B-42,306
REFERENCE/OCKET NUMBER: B-42,306
TELEPRONE: 617/542-8906
                                                                                                                                                                                                                                                                                                                                         E: Fish & Richardson P.C. 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCT-US94-02631-72; Sequence 72, Application PC/TUS9402631; GENERAL INFORMATION:
; Sequence 21, Application US/08060988A ; Patent No. 6294322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 93.3%;
Matches 14; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-060-988A-21
                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Fie
STREET: 225 Fr
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
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COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
LYMPHOCYTES AGAINST HIV
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                                                                                                                                               COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02631
FILING DATE:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
FILING DATE: 13-MAY-1994
CLASSIFICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE TITLE OF INVENTION: ELICITING NEUTRALIZING ANTITITLE OF INVENTION: IMPROCYTES AGAINST HIV NUMBER OF SEQUENCES: 36 CORRESPONDENCE ADDRESS: ADDRESSE: Birch, Stewart, Kolasch & Birch STREET: P.O. BOX 747 CITY: Falls Church STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.5%; Score 72; DB 5; Lo
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 North Figueroa Street, Suite 500
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19, Application PC/TUS9405142 GENERAL INFORMATION:
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100.0%; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TIONED TO THE TOTAL TOWN THE TOWN THE TENDENTATION NUMBER: 29,215
REGISTRATION NUMBER: 1920
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
TELEFAX: (213) 977-1001
INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIQRGPGRAFVTIG 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   : 15 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide
                                                                                                     CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
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us-08-869-386-1.rai

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TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LIMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSER: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    / Match 93.5%; Score 72; DB 5; Length 15; Local Similarity 93.3%; Pred. No. 5.2e-05; Pred. 14; Conservative 0; Mismatches 1; Indels
                                                                                                                      Score 72; DB 5; Length 15;
Pred. No. 5.2e-05;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CUNNEY: USA

ZIP: 22040-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RELEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
CLASSITCATION:
PRIOR APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: 1.715
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-13, see Table V"
                             /label= peptide
/note= "p18-12, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330 REFERENCE/DOCKET NUMBER: 1173-434P
                                                                                                                                                                                                                                                                                                                                                                             Sequence 21, Application PC/TUS9405142 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 21: SEQUENCE CHARACTERISTICS:
                                                                                                                    ch 93.5%;
11 Similarity 93.3%;
14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 RIORGPGRAFVTIGK 15
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    LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
                                                                                                                      Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                            ;
PCT-US94-05142-20
                                                                                                                                                                                                                                                                                                                                                          PCT-US94-05142-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 135
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STRET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:

SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTONER: Svensson, Leonard R.

REGISTRATION NUMBER: 30330
REGISTRATION NUMBER: 30330
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                               NAME/KEY: Peptide
LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-11, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.3%; Score 72; DB 5; 93.3%; Pred. No. 5.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8050
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acids
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20, Application PC/TUS9405142 GENERAL INFORMATION: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 20:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIQRGPGRAFYTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 133
PCT-US94-05142-20
                                                                                                                                                                                                                                                                                                                                                                                                                             PCT-US94-05142-19
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Gapa

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CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UNDER: 08/257,528
FILING DATE: 09-UNN-1994
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION WUMBER: 08/073,378
FILING DATE: 09-UNN-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION WUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-450 MIS:jb
TELEFORMUNICATION INFORMATION:
TELEFORMUNICATION INFORMATION:
TELEFORMUNICATION SEQ ID NO: 35:
SEQUENCE CHARACTERIFICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1038-487 MIS: jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10:
TELECOMMUNICATION INFORMATION:
02-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIQRGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 17 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: single
; TOPOLOGY: linear
US-08-460-602A-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Toronto STATE: Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-463-966A-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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Patent No. 5759769
GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: CHONG, Pele
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: SIM & MCBUINEY
CORRESPONDENCE ADDRESS:
                                                                                               GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: TOTORICO
STATE: Ontario
CONTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.25
SOFTWARE: PAtentin Release #1.0, Version #1.25
SOFTWARE: PAPLICATION DATA:
APPLICATION NUMBER: US/08/257,528B
FILING DATE: 09-400-1994
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAMM: STEMART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-336 MIS:jb
TELECOMMUNICATION INFORMATION:
MET DELICY FOR 115 FOR 
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ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,602A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 72; DB ]; Pred. No. 5.86 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Tozonto
STATE: Ontario
                             Sequence 35, Application US/08257528B Patent No. 5639854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93.5%; Sco.
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIORGPGRAFVTIG 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGIH: 17 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 93.5
Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: Single;
TOPOLOGY: linear
US-08-257-528B-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-460-602A-35
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Gaps
93.5%; Score 72; DB 1; Length 17; 100.0%; Pred. No. 5.8e-05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE COMPUTER: IBM PC COMPATIBLE COMPUTER: IBM PC COMPATIBLE COMPUTER: IBM PC COMPATIBLE CONTRACTOR SOCTAME: PATENTIN Release #1.0, Version #1.25 SOCTAMENT APPLICATION NUMBER: US/08/463,966A FILING DATE: 05-JUN-1995 CLASSIFICATION NUMBER: 08/257,528 FILING DATE: 09-JUN-1994 CLASSIFICATION: 424 PRIOR APPLICATION WHERE: 08/073,378 FILING DATE: 09-JUN-1994 CLASSIFICATION WHERE: 08/073,378 FILING DATE: 09-JUN-1994 CLASSIFICATION WHERE: 08/073,378 FILING DATE: 09-JUN-1993 CLASSIFICATION: 424 CLASSI
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US-08-464-329A-35
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                                                                                                                                                                                                                   Query Match 93.5%; Score 72; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 5.8e-05; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 35, Application US/08465217A

Patent No. 5800822

GENERAL INFORMATION:
APPLICANT: CHONG, Pele
TITLE OF INFORTION:
TITLE OF INFORTION:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
SIME MEDICANE
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: SULTE ''', ...

CITY: Toronto
STATE: Ontario
STATE: Ontario
STATE: Canada
ZIP: MSG IR7
COMPUTER: ELADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENT P. PC-DOS/MS-DOS
FILING DATE: O9-JUN-1995
CLASSIFICATION 1424
PRIOR APPLICATION 1424
PRIOR APPLICATION 1424
PRIOR STEMART: MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-486 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1163
TELEPHONE: (416) 595-1163
INPORMATION FOR SEQ ID NO: 35:
TELEPHONE: (416) 595-1155
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 aming acids
                                                                                                                                                                                                                                                                                                         1 RIQRGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                      4 RIORGPGRAFVTIG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17 amino acids
                                                                                                         TYPE: amino acid
STRANDEDNESS: single
; TOPOLOGY: linear
US-08-463-966A-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , TOPOLOGY:
US-08-465-217A-35
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US-08-465-217A-35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                             GITY: Sulte 'vi, ...

CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIPE MSG 1R7

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OMBUTER: IBM PC Compatible
OFERATION SYSTEM: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/464,329A
FILING DATE: 05-JUN-1995
CLASSIFICATION HOMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION NUMBER: 24,973
ATTORNEY/AGENT INFORMATION:
MANE: STEMARY NET INFORMATION:
NAME: STEMARY INFORMATION:
TELEPHONE: (416) 595-1163
INFORMATION POR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
TENETHY: 17 amino acids
Sequence 35, Application US/08464329A
Patent No. 5817754
GENERAL INFORMATION:
APPLICANT: STA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: SIM & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 35. Application US/08462507A
Patent No. 5876731
GENERAL INFORMATION:
APPLICANT: STA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: CHONG, Pele
APPLICANT: TKLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93.5%; Score 72; DB 2; Ld
100.0%; Pred. No. 5.8e-05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                   ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 RIORGPGRAFVTIG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 93.5
Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: linear
US-08-464-329A-35
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Gaps

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1 RIQRGPGRAFVTIG 14

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Gaps
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GENERAL INFORMATION:
APPLICANT: REPLIGEN CORPORATION
APPLICANT: THE ROCKEFELLER UNIVERSITY
TITLE OF INVENTION: MULTIPLE ANTIGEN PEPTIDES FOR USE AS HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93.5%; Score 72; DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: Massachusetts
COUNTRY: U.S.A.

COUNTRY: U.S.A.

COUNTRY: U.S.A.

COMPUTER: D.10-2804

COMPUTER: IBM PS/2 Model 50Z or 55SX
COMPUTER: IBM PS/2 Model 50Z or 55SX
COMPUTER: IBM PS/2 Model 50Z or 55SX
COMPUTER: D.10 PS/2 Model 50Z or 55SX
COMPUTER: IBM PS/2 MOGEL 50Z
COMPUTER: D.5 MOGEL 50Z
COMPUTER: ON MOMBER: PCT/US92/06688
FILING DATE: 19920811
CLASSIFICATION NUMBER: 744,281
FILING DATE: 13 August 1992
ATTORNEY/AGENT INFORMATION:
NAME: PRUIT CLARK
REGISTRATION NUMBER: 00231/052WO1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                      1038-488 MIS: jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                      CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: 08/073,378
PILING DATE: 09-JUN-1993
PILING DATE: 424
                                                                                                                                       FILING DATE: 0.2....
CLASSIFTATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELEPONE: (416) 595-1155
TELEPAX: (416) 595-1155
TELEPAX: (416) 595-1163
               O9-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0%; P.
Matches 14; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richarder
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (617) 542-5070
(617) 542-8906
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single;
TOPOLOGY: linear
US-08-467-881A-35
PRIOR APPLICATION DATA:
                       APPLICATION NUMBER FILING DATE: 09-JUCLASSIFICATION: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: AMINO ACID TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 142
PCT-US92-06688-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INCOMMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
                                                                                                                                                                   COMPUTER KEADABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/462,507A
FILING DATE: 05-UN-1995
CLASSIPICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-UN-1994
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION NUMBER: 1038-451 MIS:jb
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25 SOFTWARE: Patentin Release #1.0, Version #1.25 TORENT APPLICATION DATA: APPLICATION NUMBER: US/08/467,881A CLASSIFICATIONTON OG-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 72; DB 2; Le
Pred. No. 5.8e-05;
       ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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Patent No. 5951986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: Ontario
COUNTY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Rloppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 93.5%; Sc
Best Local Similarity 100.0%; P
Matches 14; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 RIORGPGRAFVTIG 17
                                                                                ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDLUM TYPE: Flower:
COMPUTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-462-507A-35
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APPLICANT: Berzofsky, Jay A.

APPLICANT: Alers, Jeffrey D.

APPLICANT: Alers, Jeffrey D.

APPLICANT: Pendleton, C. D.

APPLICANT: Pendleton, C. D.

APPLICANT: Pendleton, C. D.

APPLICANT: Shirai, Muteunorii

TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT

TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSEE: Birch, Stewart, Kolasch & Birch

STREET: P.O. Box 747

CITY: Falls Church

STATE: Virginia

COUNTY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92.2%; Score 71; DB 2; Length 15; 93.3%; Pred. No. 7.4e-05; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
FILING DATE:
PRIOR APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
ANNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 1.15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-5, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
TELEPAX: 703-205-8050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-455-625-15
Sequence 15, Application US/08455625
Patent No. 591218
GENERAL "INFORMATION:
APPLICANT: Ablers, Jeffrey D.
APPLICANT: Pendleton, C. D.
                                                            ; Sequence 13, Application US/08455625
; Patent No. 593218
GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 93.3:
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nara, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-455-625-13
                                              US-08-455-625-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE
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Sequence 11, Application US/08455625

Sequence 11, Application US/08455625

Sequence 11, Pendation:

APPLICANT: Berzofsky, Jay A.

APPLICANT: Alers, Jeffrey D.

APPLICANT: Pendlaton, C. D.

APPLICANT: Shiral, Muteunori

TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT

TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSEE: Birch, Stewart, Kolasch & Birch

STREET: P.O. Box 747

CITY: Falls Church

STATE: Virginia
                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                           Query Match 93.5%; Score 72; DB 5; Length 17; Best Local Similarity 100.0%; Pred. No. 5.8e-05; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
92.2%; Score 71; DB 2; Length 15;
Best Local Similarity 93.3%; Pred. No. 7.4e-05;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 22040-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-3, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 14 MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEFHONE: 703-205-8000
TELEFACOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
PCT-US92-06688-13
                                                                                                                                                                                                                                                                                 US-08-455-625-11
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1 RIQRGPGRAFVTIGK 15
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                                                                     Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Peptide
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COUNTRY:
            TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STREET: P.O. Box 747
CITY: Palls Church
CONTY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Decompatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SUSTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:

APPLICANT: Berzofsky, Jay A.

APPLICANT: Berzofsky, Jay A.

APPLICANT: Ablers, Jeffrey D.

APPLICANT: Pendleton, C. D.

APPLICANT: Nara, Peter

APPLICANT: OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT

TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T

TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSE: Birch, Stewart, Kolasch & Birch

STREET: P.O. Box 747

CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gapa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: 1.15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-7, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 92.2%; Score 71; DB 2; I
Best Local Similarity 93.3%; Pred. No. 7.4e-05;
Matches 14; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INPORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 16, Application US/08455625
; Patent No. 5932218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
GY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Virginia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 146
US-08-455-625-16
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CONTINE REPARABLY OF THE ACCOUNT REPARABLY OF
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US-08-455-685-11
; Sequence 11, Application US/08455685
; Patent No. 6214347
                                                                                                                                                                                                                                92.2%;
                                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                           1 RIORGPGRAFVTIAK 15
   TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                    Query Match
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM
MEDIUM TYPE: Diskett
                                                                                                                                         NAME/KEY: Peptide

// CCATION: 1..15

OTHER INFORMATION:

// OTHER INFORMATION:

US-08-455-625-22
                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Berzof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 2
CITY: BOE
STATE: MA
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APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. D.
APPLICANT: Pendleton, C. D.
APPLICANT: Pendleton, C. D.
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: LIMPHOCYTES AGAINST HIV
NUMBER OP SEQUENCES: 36
CORRESPONDENCE ADDRESS: 36
CORRESPONDENCE ADDRESS: ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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0
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                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
                                                                                                                                                                                                                                                                                                                                                                            92.2%; Scur.
93.3%; Pred. No. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Svensson, Leonard R.
RECISTRATION NUMBER: 30330
REPERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION:
TELEPHONE: 703-205-8000
                                                                                                              1173-434P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                      ns 08/060,988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22, Application US/08455625
Patent No. 593218
GENERAL INFORMATION:
1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIORGPGRAIVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 93.33
Best Local Similarity 93.33
                                                                                                                                                                                                                                             MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
FRATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
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APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Briari, Muteunori
TITLE OF INVENTION: MULTIDEFERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AN
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND BESS OCRRESPONDENCE ADDRESS:
ADDRESSEE: Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                   ö
                                                                                                          Score 71; DB 2; Length 15;
Pred. No. 7.4e-05;
0; Mismatches 1; Indels
/label= peptide
/note= "p18-14, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: Diskette
COMPUTER: Diskette
COMPUTER: IBM Comparible
OPERATING SYSTEM: Windows95
SOFFWARE: FeatSEQ for Windows Version 2.0
SURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/060,988
FILING DATE: 14-MAY-1993
FILING DATE: 06-MAY-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/751,998
FILING DATE: 25-AUG-1991
APPLICATION NUMBER: 07/7148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Beatle, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE DOCKET NUMBER: 08830/022003
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 200154
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
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1 RIORAPGRAFVTIGK 15
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: peptide US-08-455-685-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 14; Conserval
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APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Shizai, Mutsunori
TITLE OF INVENTION: MULTIDER THAT ELICIT
TITLE OF INVENTION: HELPER T.LYMPHOCYTE, CYTOTOCXIC T LYMPHOCYTE AND TITLE OF INVENTION: HELPER T.LYMPHOCYTE, CYTOTOCXIC T LYMPHOCYTE AND TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                              Gaps
                                                                                                                                                                                              ö
                                                                                                                                   Score 71; DB 3; Length 15; Pred. No. 7.4e-05;
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                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTER READABLE FORM:
MEDIUM TYPE: Diskette COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette COMPUTER: Diskette COMPUTER: Diskette COMPUTER: DISKETTE Windows95
SOFTWARE: FRSESO for Windows Version 2.0
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
FILING DATE: 08/060,988
FILING DATE: 06-MAR-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1993
APPLICATION NUMBER: 07/751,998
FILING DATE: 20-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 20-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY AGENT INFORMATION:
NAMME: BEALTIE, INGTAMATION:
NAMME: BEALTIENCE NOUNDER: 0417/542-6006
                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13, Application US/08455685
Patent No. 6214347
                                                                                                                                   92.2%;
93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 200154
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                RITRGPGRAFVTIGK 15
                                                                                                                                                                                                                                              1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 93.33
Matches 14, Conservative
                                                                                                                                                                                        14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide
                                                                                                                                Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
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                                                             7 MOLECULE 111
US-08-455-685-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-455-685-13
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                                                                                                                             APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MUTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AN
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
CITY: BOSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                                                                                                                                                                                                                                                                                                          CITY: Boston
STATE: MA
COUNTY: US
ZIP: 02110-2804

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: TBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASESE for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 14-MAY-1995
PRILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 02-AUG-1991
APPLICATION NUMBER: 07/751,998
FILING DATE: 28-JAN-1988
FILING DATE: 28-JAN-1988
ATTORNEY/AGENT INFORMATION:
ANAMERY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 71; DB 3; I
Pred. No. 7.4e-05;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08830/022003
US-08-455-685-15; Application US/08455685; Sequence 15, Application US/08455685; Patent No. 6214347; GENERL INFORMATION: APPLICANT: Ablers, Jeffrey D.; APPLICANT: Ahlers, Jeffrey D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         %208-455-685-16
; Sequence 16, Application US/08455685
; Patent No. 6214347
; GENERAL INFORMATION:
; APPLICANT: Berzofsky, Jay A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-42,306
                                                                                          Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ch 92.2%;
1 Similarity 93.3%;
14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-4: REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIORGPARAFVTIGK 15
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617/542-8906
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Gaps

us-08-869-386-1.rai

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92.2%; Score 71; DB 3; Length 15; 93.3%; Pred. No. 7.4e-05; tive 0; Mismatches 1; Indels
                                                                                                                             COUNTRY: US
ZIP: 02110-2804

COMPUTER READABLE FORM:
MEDIUM TYPE: Disfecter
COMPUTER: BM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/847,311
FILING DATE: 26-MQ-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-198
FILING DATE: 26-JAN-198
RIGIGTRATION NUMBER: P-42,306
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: 08830/022003
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 22, Application US/08455685; Patent No. 6214347
                    225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIQRGPGRAIVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 92.2
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
                                                       Boston
                                                                                                                      SD
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APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18, Application US/08455685

Patent No. 6214347
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Shiral, Muteunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES:
CORRESPONDENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible OPERATING SYSTEM: Windows95 SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08830/022003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
FILING DATE: 14-MAY-1995
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 06/66,988
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-40G-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 29-40G-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: 08030/022
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fish & Richardson P.C.
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Best Local Similarity 93.3<sup>3</sup>
Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-455-685-16
                                                                                                                                                                                                                                                                                                                                                                                           Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-455-685-18
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GENERAL INFORMATION:
APPLICANT: Berzofeky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendlecon, C. David
APPLICANT: Pendlecon, C. David
APPLICANT: Shirai, Muteunori
TITLE OF INVENTION: MUTIDEFERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES:
ADDRESSER: Fish & Richardson P.C.
STREET: 225 Franklin Street
Gaps
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APPLICANT: Berzofsky, Jay A.
APPLICANT: Ablers, Jeffrey D.
APPLICANT: Ablers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: HULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
CORRESPONDENCE ADDRESS: 48
WUMBER OF SEQUENCES 48
WUMBERSEPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                  Score 71; DB 3; Length 15;
Pred. No. 7.4e-05;
0; Mismatches 1; Indels
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ZIF: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY AGENT INFORMATION:
                                ATTORNEY/AGENT INFORMATION:
NAME: BEAETICE, INGTID A.
REGISTRATION NUMBER: P-42.306
REFERENCE/DOCKET NUMBER: 08830/022001
TELECOMMUTCATION INFORMATION:
TELEPHONE: 617/542-5070
TELEPRAK: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Fish & Richardson P.C. STREET: 225 Franklin Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 13, Application US/08060988A Patent No. 6294322 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 0.
TELECOMMUNICATION INFORMATION
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                       92.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
                                                                                                                                                            TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 15 anima acids
                26-JAN-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-4
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 93.33
Matches 14; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                    , MOLECULE TYPE: peptide US-08-060-988A-11
                                                                                                                                                                                                                                                                                    amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ns
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US-08-060-988A-13
                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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CITY: B
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US-060-988A-11
; Sequence 11. Application US/08060988A
; Patent No. 6294322
; GENERAL INFORMATION:
; APPLICANT: Beracfsky, Jay A.;
APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
; APPLICANT: Nara, Peter
; TITLE OF INVENTION: THAT ELICIT
; TITLE OF INVENTION: THAT ELICIT
; TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
; NUMBER OF SEQUENCES, 48
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
FILING DATE: 14-MAY-1993
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 06/060,988
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 05-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 25-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022003
TELEFAK: 617/542-8906
TELEFAK: 617/542-5070
TELEFAK: 200154
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 71; DB 3
Pred. No. 7.4e-
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEB: Fish & Richardson P.C. : 225 Franklin Street Boston
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APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-11992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92.2%;
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Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide US-08-455-685-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
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Conservative
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ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
les 14; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                   Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
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Matches
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                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A.
APPLICANT: Anlers, Jeffrey D.
APPLICANT: Anlers, Jeffrey D.
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
ITLE OF INVENTION: THAT ELICIT
ITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
ITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92.2%; Score 71; DB 3; Length 15; 93.3%; Pred. No. 7.4e-05; tive 0; Mismatches 1; Indels
                                                                                                               Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: DISKELLE
COMPUTER: IBM COMPALIALIE
OPERATING SYSTEM: Windows95
SOFTWARE: FASESG for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-WAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 05-WAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: BEGLISE, INGIA A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022001
TELEPHONE: 617/542-8906
TELEX: 200154
TELEX: 200154
TELEX: 200154
TELEX: 200154
TELEX: 200154
                                                                                                             92.2%; Score 71; DB 3; I 93.3%; Pred. No. 7.4e-05; Live 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                    Sequence 15, Application US/08060988A Patent No. 6294322
                                                                                                                                                                                              1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                 1 RIORAPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 93.33
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           : 15 amino acids
amino acid
15 amino acids
                                                                                                                                  Best Local Similarity 93.3
Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                            TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide
                amino acid
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                                                                          US-08-060-988A-13
                                                                                                                                                                                                                                                                                               RESULT 157
US-08-060-988A-15
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                                                                                                                 Query Match
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Sequence 16, Application US/08060988A

Sequence 16, Application US/08060988A

Petent No. 6294322

GENERAL INFORMATION:

APPLICANT: Berzofsky, Jay A.

APPLICANT: Ahlers, Jeffrey D.

APPLICANT: Nara, Peter

APPLICANT: Nara, Peter

APPLICANT: Nara, Peter

TITLE OF INVENTION: THAT ELLCIT

TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE.

TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE.AND

TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE.AND

TITLE OF INVENTION: HELPER T-LYMPHOCYTE, AND

TITLE OF INVENTION: HELP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Pred. No. 7.4e-05;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING APPLICATION DATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
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REGISTRATION UNBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022001
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Fish & Richardson P.C.
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18, Application US/08060988A Patent No. 6294322 GENERAL INFORMATION: APPLICANT: Berzofsky, Jay A. APPLICANT: Ahlers, Jeffrey D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELBEAK: 617/542 TELBEAK: 200154
INFORMATION FOR SEQ ID NO: 16
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92.2%;
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Gaps ; 0

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FILING DATE: 13-MAY-1994
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 08
TELECOMMUNICATION:
                                                                                                                                          COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  : 617/542-5070
617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IRELEX. 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                          COMPUTER READABLE FORM MEDIUM TYPE: Disket
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: C
TITLE OF INVENTION: E
TITLE OF INVENTION: L
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 14; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
                                                  Y: US
02110-2804
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TELEFAX: 61
TELEX: 2001
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                                                              TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELPER T.LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELPER T.LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBERS OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MULTIDETERMINANT PEPTIDES
THAT ELICIT
HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 15;
                                                                                                                                                                                                                                                                                                        COMPUTER 105

ZIP: 02110-2804

COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEC for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 03/08/060,988A
FILING DATE: 14-MAY 1993
PRIOR APPLICATION DATA:
FILING DATE: 05-MAR.1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 25-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 25-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 25-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 15-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 15-MG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 15-MG-1991
APPLICATION NUMBER: 07/148,692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7.4e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-42,306 REFERENCE/DOCKET NUMBER: 08830/022001
                                                                                                                                                                                                                E: Fish & Richardson P.C.
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Fish & Richardson P.C.
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22, Application US/08060988A Patent No. 6294322 GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A.
Pendleton, C. David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                               Shirai, Mutsunori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 93.3%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIORGPGRAIVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                  Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-060-988A-18
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US-08-060-988A-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
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COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC
LYMPHOCYTES AGAINST HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSE: Birch, Stewart, Kolasch & Birch STREET: P.O. Box 747
STREET: P.O. Box 747
CITY: Falls Church STATE: Virginia COWTRY: USA
CONFORTE: Bish Spanner: USA
COMPUTER READABLE FORM: WEDIUM TYPE: ISlopy disk
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
SOGTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
RAPPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
UPERATING SYSTEM: "Windows SOFTWARE: PastSRQ for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION UNDER: 07/847,311
                                                                                                                                                                  APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
ATONNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11, Application PC/TUS9405142
GENERAL INFORMATION:
APPLICANT:
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TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE BITCH, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STRATE: Virginia
COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                   92.2%; Score 71; DB 5; Length 15; 93.3%; Pred. No. 7.4e-05; cive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE PATENTIN STILL SOFTWARE WITH SOFTWARE SOFTWARE PATENTIN BELEASE #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142 FILING DATE: 13-MAY-1994 CLASSIFICATION DATE: PRIOR APPLICATION NUMBER: US 08/060,988 FILING DATE: 14-MAY-1993 ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330 REFERENCE/DOCKET NUMBER: 1173-434P
                        /label= peptide
/note= "p18-5, see Table V"
                                                                                                                                                                                                                                                                                                                                                       PCT-US94-05142-15; Sequence 15, Application PC/TUS9405142; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 15: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 RIORGPGRAFVTIGK 15
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                                                                                                                                                                    14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide FRAGMENT TYPE: internal
  LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 14; Conserv
                                                                                                                                              Best Local Similarity
                                                                       PCT-US94-05142-13
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Sequence 13, Application PC/TUS9405142
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
INTIE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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SOFTWARE PATENTIN Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US94/05142

FILING DATE: 13-MAY-1994

CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/060,988

FILING DATE: 14-MAY-1993

ATTORNEY/AGENT INFORMATION:

NAME: Svensson, Leonard R.

REGISTRATION NUMBER: 30330

REGISTRATION NUMBER: 30330

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEPERAS: 703-205-8050
                                                                                                                                                                                                                                                                                                                                NAME/KEY: Peptide
LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-3, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92.2%; Score 71; DB 5; I 93.3%; Pred. No. 7.4e-05; tive 0; Mismatches 1;
NAME: Svenseon, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
TELEPHONE: 703-205-8050
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acids
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 22040-0747
COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 RITRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14; Conservative
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FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                PCT-US94-05142-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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0; Gaps ch 92.2%; Score 71; DB 5; Length 15; 1 Similarity 93.3%; Pred. No. 7.4e-05; 14; Conservative 0; Mismatches 1; Indel8 LOCATION: 1..15 OTHER INFORMATION: /label= peptide OTHER INFORMATION: /note="pl8-7, see Table V"

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RESULT 164

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FILING DATE: 13-MAY-1994
FILING DATE: 13-MAY-1994
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: SYGNESON, LEGNARG R.
REGISTRATION NUMBER: 30330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCT-US94-05142-22; Sequence 22, Application PC/TUS9405142; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 703-205-8050
INPORANTION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   92.2%;
93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 93.3*
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: 1.15
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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GENERAL INFORMATION:
APPLICANT:
APPLICANT:
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: BLICTTING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 3
CORRESPONDENCES: ADDRESSEE: Blich, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Palls Church
STREET: VISAIRS
COUNTRY: USA
                                                                            TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                            COUNTRY: USA

ZIP: 22040-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-8, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    yeary match
92.2%; Score 71; DB 5; I
Best Local Similarity 93.3%; Pred. No. 7.4e-05;
Matches 14; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
         ; Sequence 16, Application PC/TUS9405142; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: Peptide
LOCATION: 1..15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22040-0747
PCT-US94-05142-16
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APPLICANT:
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 71; DB 5; Length 15;
Pred. No. 7.4e-05;
0; Mismatches 1; Indels
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: TBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWAREN PAPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
FLING DATE: 13-MAY-1994
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: SVENBSON, LEGNARD R.
REGISTRATION NUMBER: 30330
REFERENCE/POCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= peptide
/note= "p18-10, see Table V"
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Gaps

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APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Bendleton, C. David
APPLICANT: Shirai, Musunori
APPLICANT: Shirai, Musunori
APPLICANT: Shirai, Musunori
APPLICANT: Convexion: Multideterminant Preprides That Elicit
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AN
ATITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89.6%; Score 69; DB 3; Length 15; 93.3%; Pred. No. 0.00015;
                                                                                                                                                                         89.6%; Score 69; DB 2; Length 15; 93.3%; Pred. No. 0.00015; tive 0; Mismatches 1; Indels
                                                                                     /label= peptide
/note= "p18-6, see Table V"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FaseSES for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 13-MAY-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/060,988
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 05-MAR-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: BEALLIS, INGTIA A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022003
TELEPHONE: 617/542-5070
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Fish & Richardson P.C. 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 14, Application US/08455685 Patent No. 6214347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 617/572 TELEX: 200154
INFORMATION FOR SEQ ID NO: 14: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                   1 RIQRGAGRAFVTICK 15
                                                                                                                                                                             Query Match 89.6
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide US-08-455-685-14
                                                             LOCATION: 1.15
OTHER INFORMATION:
                                          NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: MU
COUNTRY:
                                                                                                                                   US-08-455-625-14
                                                                                                                                                                                                                                                                                                                                                                                                         US-08-455-685-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 14, Application US/08455625
Patent No. 593218
GENERAL INFORMATION:
APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffray D.
APPLICANT: Pendlaton, C. D.
APPLICANT: Shiral, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                           Score 71; DB 5; Length 15;
Pred. No. 7.4e-05;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOCTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/455,625

FLING DATE:

CLASSIFICATION: 435

PRIOR APPLICATION NUMBER: US 08/060,988

ATHORISY/AGENT INFORMATION:

APPLICATION NUMBER: US 08/060,988

FLING DATE: 14-MAY-1993

ATHORISY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                      ADDRESSEE: Birch, Stewart, Kolasch & Birch STRENT: P.O. Box 747
CITY: Falls Church STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Svenson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION:
TELEPHONE: 703-205-8000
REFERENCE/DOCKET NUMBER: 1173-434P
              TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
TELEPHONE: 703-205-8050
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 93.3%;
Matches 14; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIORGPGRAFVTIAK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: peptide
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XX: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: 1.15
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCT-US94-05142-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 171
US-08-279-906A-17
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                                                                                                                                                                                                                                                                            APPLICANT: BEIZOSISKY, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. David
APPLICANT: Pendleton, C. David
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CORRESPONDENCE ADDRESS: 48
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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  Indels
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    1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 07/847,311
FILING DATE: 06-WAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A:
REGISTRATION NUMBER: 042,306
REFERENCE/DOCKET NUMBER: 08830/022001
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 0.00
0; Mismatches
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                                                                                                                                                                                     US-08-060-988A-14
; Sequence 14, Application US/08060988A
; Patent No. 6294322
; GENERAL INFORMATION;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Fish & Richardso
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 14:
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                                                                                         1 RIORGAGRAFVTIGK 15
                                               1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIORGAGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 93.33
Matches 14; Conservative
14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 170
PCT-US94-05142-14
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Matches
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TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSER: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STREET: Vigaina
COUNTRY: USA
ZIP: 22040-0747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Patent No. 5618922
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Terada, Masaki
APPLICANT: Yoneda, Yukio
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray &
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray &
STREET: 6300 Sears Tower, 233 S. Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC COMPAGE OF COMPAGE OF STREM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /label= peptide
/note= "p18-6, see Table V"
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STREET: 6300 Sears Tower, 233 S. Wacker Drive
CITY: Chicago
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 69; DB 5; ]
Pred. No. 0.00015;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 13-MAY-1994
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   89.6%;
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Gaps

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APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleron, C. David
APPLICANT: Pendleron, C. David
APPLICANT: Nara, Peter
APPLICANT: Shira, Mitsunori
TITLE OF INVENTION: MILTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AN
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                       Length 14;
                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                        NAME/KEY: Peptide
| NAME/KEY: Peptide
| LOCATION: 1..14
| OTHER INFORMATION: /label= peptide
| US-08-455-625-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastENG for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/45,685
FILING DATE: 31-MAY-1995
PRION APPLICATION NUMBER: 08/060,988
FILING DATE: 4-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY-AGENT INFORMATION:
NAME: BEAELIE, INGILA A.
NAME: BEAELIE, INGILA A.
NAME: BEAELIE, INGILA A.
                                                                                                                                                                                                                                                                                                                                    Score 68; DB 2;
Pred. No. 0.0002;
                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08830/022003
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225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10, Application US/08455685
Patent No. 6214347
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                 88.3%; Sc.
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 08:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEBRAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          :0
              10:
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STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                               3 QRGPGRAFVTIGK 15
                                                         14 amino acids
                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 ORGPGRAFVTICK 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 14 amino acids
              INFORMATION FOR SEQ ID NO:
                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
STREET: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-455-685-10
                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. D.
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STRIE: Virgina
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/279,906A
PILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PPRIOR APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REGISTRATION NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                    CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: No. 5618922and, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 32028
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10, Application US/08455625
Patent No. 5932218
                                                                                                                                                                                                                                                                                                             TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-6448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 QRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGPGRAFVTIGK 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-279-906A-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 172
US-08-455-625-10
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LOCATION: 1.14
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "pl8-2, see Table V"
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Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: SYCHERON, LECOLARD R. REGISTRATION NUMBER: 30330 REFERENCE/DOCKET NUMBER: 1173-434P TELECOMMUNICATION INFORMATION: 703-205-8000 TELEFAX: 703-205-8050
                                                                               Sequence 10, Application PC/TUS9405142 GENERAL INFORMATION: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 51, Application US/08257528B ; Patent No. 5639854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 ORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 ORGPGRAFVTIGK 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
                                          RESULT 175
PCT-US94-05142-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCT-US94-05142-10
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US-08-257-528B-51
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                                                                                                                                                                                                                                                                                                                      Sequence 10, Application US/0806098BA

Sequence 10, Application US/0806098BA

Patent No. 6294322

GENERAL INFORMATION

APPLICANT: Berzofsky, Jay A.

APPLICANT: Berzofsky, Jay A.

APPLICANT: Anlers, Jeffrey D.

APPLICANT: Nara, Peter

APPLICANT: Nara, Peter

TITLE OF INVENTION: WILTIDETERMINANT PEPTIDES

TITLE OF INVENTION: THAT ELICIT LYMPHOCYTE AND

TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1

NUMBER OF SEQUENCES: 48

CORRESPONDENCE ADDRESS:

ADDRESSE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                      Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: DISKETE COMPUTER: JEBM COMPAGE COMPUTER: LIBM COMPAGE CONTROL OF WINDOWS SOFTWARE: FASEISED for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A FILING DATE: 14-MAY-1993
PRIOR APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 26-JAN-198
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-198
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-198
ATTORNEY/AGENT INFORMATION:
NAME: BEATLE, INGIT A.
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: P-42,306
REGISTRATION NUMBER: P-42,306
TELECHOMMUNICATION INFORMATION:
TELEPHONE: 617/642-5070
                                                                                    Query Match 88.3%; Score 68; DB 3; 18est Local Similarity 100.0%; Pred. No. 0.0002; Matches 13; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Fish & Richardson P.C.
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 200154
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                3 ORGPGRAFVTIGK 15
                                                                                                                                                                                                         2 QRGPGRAFVTIGK 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 ORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGPGRAFVTIGK 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diskette
                     , MOLECULE TYPE: peptide US-08-455-685-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Boston
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                                                                                                                                                                                                                                                                                                              US-08-060-988A-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
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TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STREET: Vigilia
COUNTRY: USA
ZIP: 22040-0747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Plei
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: SIM & MCBUINEY
                                                                                                                                                                                                                                                                                      ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
CLASSIFICATION SATE: 08/060,988
FILING DATE: 14-MAY-193
ATYORNEY/AGBNT INPORMATION:
NAMP: CLASSIFICATION TOWNER: US 08/060,988
FILING DATE: 14-MAY-193
ATYORNEY/AGBNT INPORMATION:
NAMP: CLASSIFICATION TOWNER: US 08/060,988
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Score 67; DB 1; Length 20; Pred. No. 0.00041;
                                                                                                                                                                                                                                                                                                                                             87.0%; Score 67; DB 1; I
92.9%; Pred. No. 0.00041;
tive 0; Mismatches 1;
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1163
TELEPHONE: (416) 595-1163
INFORMATION FOR SEQ ID NO: 51: SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: amino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              87.0%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIQRGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7 RIORGPGRAFYTIG 20
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                              ; TOPOLOGY: linear
US-08-460-602A-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-463-966A-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-463-966A-51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        ઠે
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US-08-460-662A-51
i Sequence 51, Application US/08460602A
j Patent No. 5759769
j GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,602A
PILING DATE: 09-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY AGENT INFORMATION:
NAME: STEMART, MICHAEL I.
                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/257,528B
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                             1038-336 MIS: jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPAX: (416) 595-1163
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RIORGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7 RIORGPGRAFYTIG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13; Conservative
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Best Local Similarity
Matches 13; Conserva
                              8: Ontario
RY: Canada
MSG 1R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-257-528B-51
                                                         COUNTRY:
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Gaps
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Patent No. 5795955
GENERAL INFORMATION:
APPLICANT: CHONG. Pele
APPLICANT: CHONG. Pele
APPLICANT: KIEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
CORRESPONDENCE ADDRESS: 101
CORRESPONDENCE ADDRESS: 300
CORNTRY: Canada
ZIP: MGG 1R7
COMPUTER: Suite 701, 330 University Avenue
CITY: Toronto
COMPUTRY: Canada
ZIP: MGG 1R7
COMPUTRY: IBM PC Compatible
COMPUTRY: BEADABLE FORM:
MEDILOMITYPE: FLOPY disk
COMPUTRY: BEADABLE FORM:
APPLICATION TYPE: GAS-IN-1995
CLASSIFTCATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFTCATION: 424
PRIOR APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1994
CLASSIFTCATION: 424
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 24,973
REFERENCE/ACTION ACTION NUMBE
Length 20;
                                                                                    1; Indels
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87.0%; Score 67; DB 2; Length 20; 92.9%; Pred. No. 0.00041; ive 0; Mismatches 1; Indels
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US-08-462-507A-51
Sequence 51, Application US/08462507A
Sequence 51, Application US/08462507A
Setent No. 5876731
GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Sulte 701, 330 University Avenue
STARET: Ontario
CITY: Toronto
STARET: Ontario
COUNTRY: Canada
SIRE MSG 1R7
SI
                                                                                                                                                                                                                                                                       COMPUTER: RELORDY disk
COMPUTER: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPANIOS SYSTEM: PC-DOS/MS-DOS
SOFFWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,329A
FILING DATE: 05-JUN-1995
CLASSIFICATION NUMBER: 08/257,528
FILING APPLICATION 1424
PRIOR APPLICATION: 424
CLASSIFICATION: 424
CLASSIFICATION: 424
CLASSIFICATION: 424
CLASSIFICATION: 424
CLASSIFICATION: 424
CLASSIFICATION: 424
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
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                          NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MGG IR7
COMPUTER READABLE
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,913
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEFRAN: (416) 595-1155
TELEFRAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PC compatible
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 13; Conserva
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; TOPOLOGY: linear
US-08-464-329A-51
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  Indels
                                                                                                                                                                                                                                    Sequence 51, Application US/08465217A

Patent No. 5800822

GENERAL INFORMATION:

APPLICANT: SIA, Charles D.Y.
APPLICANT: KLEIN, Michel H.

TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101

CORRESPONDENCE ADDRESS:

ADDRESSEE: Sim & McBurney

STREET: Suite 701, 330 University Avenue

CITY: Toronto

STATE Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 180
20.08-464-329A-51
; Sequence 51, Application US/08464329A
; Patent No. 5817754
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APPLICANT: SIA, Charles D.Y.
APPLICANT: CHOWG, Pele
APPLICANT: KLEIN, Michel H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RIQRGPGRAFVTIG 14
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                                                             1 RIQRGPGRAFVTIG 14
                                                                                                                  7 RIORGPGRAFYTIG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20 amino acids
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        13; Conservative
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linear
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY:
US-08-465-217A-51
                                                                                                                                                                                                                                 US-08-465-217A-51
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        Matches
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0; Gaps

Gaps

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APPLICANT: CARPIO MUSCZ, E. L. APPLICANT: CARPIO MUSCZ, E. L. APPLICANT: QUINTANA V. D. APPLICANT: GIVEZ CODR GUEZ, C. E. APPLICANT: SILVA RODR GUEZ, R. C. APPLICANT: SILVA RODR GUEZ, R. C. APPLICANT: NAZ BAL GIVEZ, C. APPLICANT: ILEAL ANGULO, M. J. TITLE OF INVENTION: System for the expression of heterologous TITLE OF INVENTION: antigens as fusion proteins CORRESPONDENCE ADDRESS: 21
CORRESPONDENCE ADDRESS: Lackenbach Siegel Marzullo Aronson & Greenspan
                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Lackenbach Siegel Marzullo Aronson & Greenspan STREET: One Chase Road CITY: Scarsdale STAIE: New York COUNTRY: U.S.
                                                                                                                                                                                                                                                                                                                                                   Score 67; DB 2; Length 20;
Pred. No. 0.00041;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk 3.5'' (1.4 MB).
COMPUTER: Compatible PC IBM (80486, 8 M Ram).
COMPUTER: Windows 95.
SOFTWARE: Word Perfect 5.0 for Windows 95.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/930,917A
FILING DATE: 15-Sep-1997
CLASSIFICATION NUMBER: PCT/CU97/00001
FILING DATE: 17-Jan-1997
ATTORNEY/AGENT INFORMATION:
NAME: HENRY A. MARZULLO, JR.
REGISTRATION NUMBER: 20,910
RESTRENCE/DOCKET NUMBER: P-13
TELECOMMUNICATION INFORMATION:
TELEFHONE: (914) 723-4300
TELEFHONE: (914) 723-4300
TELEFRONE CHARACTERISTICS:
LEMGTH: A.15 AMINO acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14, Application US/08930917A; Patent No. 6146635; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DUARTE CANO, C. A. GUILL N NIETO, G. E MART N DUNN, A. M. ALVAREZ ACOSTA, A.
   TELECOMMUNICATION INFORMATION: TELEPHONE: (416) 595-1155
                                                         TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 51
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                             87.0%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 RIORGEGRAFYTIG 20
                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 92.9
Matches 13; Conservative
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STRANDEDNESS: Unkr
TOPOLOGY: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ANTI-SENSE: No
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                                                                                                                                                                                                                                                                                             US-08-467-881A-51
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Pred. No. 0.00041;
0; Mismatches 1; Indels
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APPLICATION NUMBER: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,881A
FILING DATE: 06-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
PRIOR APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
PRIOR APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
PRIOR APPLICATION: 424
                    FILING DATE: 05-UW-1995
CLASSIFICATION: 424
PRIOR APPLICATION 1424
PRIOR APPLICATION 1424
PRIOR APPLICATION 1424
PRIOR APPLICATION 424
PRIOR APPLICATION 424
PRIOR APPLICATION 424
PRIOR APPLICATION 1993
CLASSIFICATION 1993
CLASSIFICATION 1993
CLASSIFICATION 1993
CLASSIFICATION 1993
FILING DATE: 09-UN-1993
CLASSIFICATION 100-1993
FILING APPLICATION 100-1993
CLASSIFICATION 100-1993
FILING DATE: 09-UN-1993
CLASSIFICATION 100-1993
FILING AREA 100-1993
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REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-488 MIS:jb
APPLICATION NUMBER: US/08/462,507A FILING DATE: 05-JUN-1995 CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   87.0%;
92.9%;
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Best Local Similarity 92.9
Matches 13; Conservative
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linear
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13 RIQRGPGRAFVTI 25

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amino acid
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US-08-279-906A-19
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US-08-704-170-52
                                                                                                                                                                                                                                                                                                                                                    CITY: Ch:
STATE: I.
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX:
TELEX: 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Svennerholm, Bo
APPLICANT: Svennerholm, Bo
APPLICANT: Svennerholm, Bo
APPLICANT: Genasson, Stig
APPLICANT: Horal, Peter
TITLE OF INVENTION: INDUCTION OF NEUTRALIZING ANTIBODIES AGAINST HUMAN
TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS
NUMBER OF SEQUENCES: 41
CORRESPONDENCES: 41
CORRESPONDENCES: ADDRESS:
ADDRESSEE: KNOBBE, MARTENS, OLSON AND BEAR
STREET: 620 NEWPORT CENTER DRIVE 16TH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                Gaps
                                                                                                Central region of the loop V3 belonging to the protein gpl20 from the VIH-1, isolation IIIB.
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85.7%; Score 66; DB 2; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.00074;
Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                      85.7%; Score 66; DB 3; Length 15; 100.0%; Pred. No. 0.00045; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE:
COUNTRY: USA
ZIP: 92660
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/493,235
FILING DATE: 20) June)1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Kaiser, Annewarie
REGISTRATION NUMBER: 37,649
REGISTRATION NUMBER: 37,649
REGISTRATION NUMBER: 37,649
REGISTRATION INFORMATION:
TELECHONE: 619-235-8550
TTELEPHONE: 619-235-8550
TTELEPHONE: 619-235-8550
TTELEPHONE: TSEQ ID 00: 23:
                                                                                                                                                                                                                                                                                                                                                                                                    US-08-493-235-23
; Sequence 23, Application US/08493235
; Patent No. 5840313
FRAGMENT TYPE: Internal fragment
ONGINAL SOURCE:
ORGANISM: VIH-1
INDIVIDUAL ISOLATE: IIIB
                                                                                                                                                                                                                                                                            1 RIQRGPGRAFVTI 13
                                                                                                                                                                                                                                                                                                      3 RIQRGPGRAFVTI 15
                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RAGMENT TYPE: internal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                       OTHER INFORMATION:
COTHER INFORMATION:
US-08-930-917A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ANTI-SENSE:
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1 RIORGPGRAFVII 13

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US-08-278-906-1.

US-08-278-906-1.

US-08-278-906-1.

Sequence No. 55.1852.

Patent No. 50.1852.

Patent No. 50.1852.

ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ATDRESSEE: Marchall. O'Toole, D'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Murray & ADDRESSEE: Marchall. O'Toole, Geretein, Geretein
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                 TELEX: 421792
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                          Query Match 81.8
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 RGPGRAFVTIGK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 RGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide
PCT-US94-02631-52
                                                                                                                                                                                                                                                         TYPE: Amino acida
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: Unknown
US-08-488-252-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 90012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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Patent No. 5763160
GENERAL INFORMATION:
APPLICANT: Chang Y1 Wang
TITLE OF INVENTION: SYNTHETIC PEPTIDES AND PROCESS
TITLE OF INVENTION: OF USING SAME FOR THE DETECTION OF ANTIBODIES TO
TITLE OF INVENTION: HOMAN INMUNOBFICIENCY VIRUS (HIV) GP120 ENVELOPE
TITLE OF INVENTION: AND AS VACCINES
TITLE OF INVENTION: AND AS VACCINES
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81.8%; Score 63; DB 1; Length 12; 100.0%; Pred. No. 0.0011; tive 0; Mismatches 0; Indels
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,170
FILING DATE: US/08/704,170
FILING DATE: 11-MAR-1993
PFILING DATE: 11-MAR-1993
PFILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRA: (213) 977-1001
TELEFRA: (213) 977-1001
TELEFRA: (213) 977-1003
TELEFRA: (213) 977-1003
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURBENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,252
FILING DATE: 07-Jun-1995
APPLICATION NUMBER: 08/326,676
FILING DATE: 07-Jun-1995
APPLICATION NUMBER: 07/26,605
FILING DATE: 09-July-1991
APPLICATION NUMBER: 07/125,321
FILING DATE: 01-MR-1991
APPLICATION NUMBER: 07/155,321
FILING DATE: 12-Feb-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : 12 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 RGPGRAFVTIGK 15
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CITY: NEW YORK
STATE: NEW YORK
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-488-252-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-704-170-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
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Score 62.5; DB 1; Length 21;
Pred. No. 0.0022;
0; Mismatches 0; Indels
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COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPANIOS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/709,209
FILING DATE: 12-AUG-1996
CLASSIFICATION 1424
PRIOR APPLICATION NUMBER: US 08/105,483
FILING DATE: 12-AUG-1993
APPLICATION NUMBER: US 07/847,951
FILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
   CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/105,483
FILING DATE: US/08/105,483
FILING DATE: US/08/105,483
CLASSIFICATION NUMBER: US/08/105,483
FILING APPLICATION DATA:
APPLICATION NUMBER: US 07/847,951
FILING DATE: 06-MAR-1992
ATTONNEY/AGENT INFORMATION:
NAME: FYCOMMER, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 25,506
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF SEQUENCE: (212) 840-0712
INFORMATION FOR SEQ ID NO: 3844:
SEQUENCE CHRARACTERISTICS:
LENGTH: 21 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 384, Application US/08709209
Patent No. 5762938
GENERAL INFORMATION:
APPLICANT: Paoletti, Enzo
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
TITLE OF INVENTION: STRAIN
NUMBER OF SEQUENCES: 462
CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris & Safford
ADDRESSEE: Cortis, Morris & Safford
ADDRESSEE: Cov #111am S. Frommer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Frommer, William S.
REGISTRATION NUMBER: 25.506
REFERENCE/DOCKET NUMBER: 454310-2400
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 530 Fifth Avenue CITY: New York STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 93.33
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
;
TOPOLOGY: linear
US-08-105-483-384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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US-08-709-209-384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 12;
                                                                                                                                                                                          Sequence 43, Application PC/TUS9503236
GENERAL INFORMATION:
APPLICANT: University of Southern California
TITLE OF INVENTION: Methods to Diagnose and Treat HIV-1
TITLE OF INVENTION: Infection
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | Sequence 384, Application US/08105483
| Sequence 384, Application US/08105483
| Patent No. 5494807
| GENERAL INFORMATION:
| APPLICANT: Paoletti, Enzo
| TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
| TITLE OF INVENTION: STRAIN
| NUMBER OF SEQUENCES: 462
| CORRESPONDERS: ADDRESSE: ADDRESSEE: Curtis, Morris & Safford
| ADDRESSEE: Curtis, Morris & Safford
| STREET: 530 Fifth Avenue
| STREET: NY OF STREET: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 92122
CMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/03236
                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRENT AFFLICATION DATA:

CARSTELATION NUMBER: PCT/US95/03236
FILING DATE: 13-MAR-1995
CLASSTELATION:
ATTORNEY/AGENT INFORMATION:
NAME: Imbra, Richard J.
REGISTRATION NUMBER: PP-SI 1394
REFERENCE/DOCKET NUMBER: PP-SI 1394
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9001
INFORMATION FOR SEQ ID NO: 43:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              81.8%; Scc. 100.0%; Pre 0;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 RGPGRAFVTIGK 15
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RGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12; Conservative
                                                      1 RGPGRAFVTIGK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
XX: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 12; Conserv
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10036
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US-08-105-483-384
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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Gaps

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Paoletti, Enzo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 194
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                                                                                                        81.2%; Score 62.5; DB 1; Length 21; 93.3%; Pred. No. 0.0022; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 62.5; DB 1; Length 21;
Pred. No. 0.0022;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                           Sequence 79, Application US/08303275
Patent No. 576559
GENERAL INFORMATION:
APPLICANT: Tartaglia, James
APPLICANT: Cox, William I.
TITLE OF INVENTION: IMMUNOBFICIENCY VIRUS RECOMBINANT
TITLE OF INVENTION: POXVIRUS VACCINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/303,275
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UNBER: US 07/897,382
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION UNBER: 25,506
REFERENCE/DOCKET NUMBER: 45,4310-2420
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Curtis, Morris & Safford ADDRESSEE: c/o William S. Frommer STREET: 530 Flfth Avenue CITY: New York STATE: New York COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 193
US-08-458-101-384
; Sequence 384, Application US/08458101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECURIO. (212) 0.0 TELEPHONE: (212) 840-0712; INFORMATION FOR SEQ ID NO: 79: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                 1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                        8 RIQRGPGRAFVT-GK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RIORGPGRAFVT-GK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 93.34
Matches 14; Conservative
21 amino acids
                                                                                                                         Best Local Similarity 93.3
Matches 14; Conservative
                 TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single
                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-08-303-275-79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: Bin
                                                       i
US-08-709-209-384
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                                                                                                                                                                                                                                                                            RESULT 192
US-08-303-275-79
                                                                                                            Query Match
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APPLICANT: Bridon, D.P.
APPLICANT: Bridon, D.P.
APPLICANT: Bridon, D.P.
APPLICANT: Bridon, D.P.
APPLICANT: Arafft, K.D.
APPLICANT: Mandecki, W.
APPLICANT: March, S.C.
APPLICANT: March, S.C.
APPLICANT: Wasell, J.R.
APPLICANT: TILE OF INVENTION: Genetically Engineered Enzymes And Their TILE OF INVENTION: Conjugates For Diagnostic Assays
NUMBER OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LABORATORIES
STREET: One Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               80.5%; Score 62; DB 2; Length 13; 100.0%; Pred. No. 0.0017; ive 0; Mismatches 0; Indels
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: SOftPC
CURRENT APPLICATION DATA:
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/100,708
FILING DATE: July 29, 1993
ATTONEY, APPLICATION:
NAME: WONG, WEAN KINING
REFRERENCE/DOCKER NUMBER: 33,561
REFRENCE/DOCKER NUMBER: 33,561
REFRENCE/DOCKER NUMBER: 33,561
TELECOMMUNICATION:
TELECOM
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/657,392
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21, Application US/08657392
Patent No. 5843634
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acid residues
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (708) 938-2623
TELEX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIQRGPGRAFVT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: unknown
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Abbott Park
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-657-392-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-657-392-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ઠે
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| Sequence 20, Application US/08657392 |
| Patent No. 5843634 |
| GENERAL INFORMATION: |
| APPLICANT: Brate, E.M. |
| APPLICANT: Bridan, D.P. |
| APPLICANT: Mandecki, W. |
| APPLICANT: Mandecki, W. |
| APPLICANT: March, S.C. |
| MUMBER OF SEQUENCES: 3 |
| CORRESPONDENCE ADDRESS: |
| AUDRESSEE: ADDRESSE: ADDRESSE: |
| ADDRESSEE: ADDRESSE: ADDRESSE: |
| CORRESPONDENCE ADDRESSE: |
| CORRESPONDENCE ADDRESSE: |
| CORRESPONDENCE ADDRESSE: |
| APPLICANT: March, Darke, Darke, March, Darke, March, Darke, March, Darke, Darke, March, Darke, Darke, March, Darke, Darke, Darke, March, Darke, Darke, Darke, March, Darke, Darke, March, Darke, Darke, March, Darke, March, Darke, Darke, March, Darke, Darke, March, Darke, Mar
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                      Genetically Engineered Enzymes And Their Conjugates For Diagnostic Assays
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/100,708
FILING DATE: July 29, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Wong, Wean Khing
REGISTRATION NUMBER: 33,561
REFERENCE/DOCKET NUMBER: 5324.US.P1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: SOFEPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/657,392
FILING DATE:
                 TITLE OF INVENTION: Genetically Eng
TITLE OF INVENTION: Conjugates For
NUMBER OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LABORATORIES
STREET: One Abbott Park Road
CITY: Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC COmpatible OPERATING SYSTEM: PC-DOS/MS-DOS
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SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acid residues
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (708) 938-2623
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MOLECULE TYPE: peptide
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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STATE: Illinois
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: USA
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US-08-657-392-20
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Length 13;
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TITLE OF INVENTION: Genetically Engineered Enzymes
TITLE OF INVENTION: And Their
TITLE OF INVENTION: Conjugates For Diagnostic Assays
NUMBER OF SEQUENCES: 34
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STATE: 111ino1s
CONTY: 0.54
CONFOUR: 0.54
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: SOFTPA
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GENERAL INFORMATION:
                                                            INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acid residues
TYPE: amino acid sTRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LABORATORIES
STREET: One Abbott Park Road
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SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: WOIG, Wean Khing
REGISTRATION NUMBER: 33.561
REFERENCE/DOCKET NUMBER: 5324
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 938-2623
TELEPHONE: (708)
                 TELEFAX: (708) 938-2623
TELEX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brennan, C.A.
Bridon, D.P.
                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                 1 RIORGPGRAFVT 12
                                                                                                                                                                     TOPOLOGY: unknown
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: unknown
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
  (108)
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                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 12; Conserv
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CLASSIFICATION:
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APPLICANT: Brate, E.M.
APPLICANT: Brate, E.M.
APPLICANT: Bridon, D.P.
APPLICANT: Bridon, D.P.
APPLICANT: Krafft, G.A.
APPLICANT: March, S.C.
APPLICANT: March, S.C.
APPLICANT: Russell, J.R.
APPLICANT: Russell, J.R.
APPLICANT: Russell, J.R.
APPLICANT: Russell, J.R.
APPLICANT: Nussrinon: Conjugates For Diagnostic Assays
NUMBER OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
ADDRESSEB: ABBOT LABORATORIES
STREET: One Abbott Park Road
CITY: Abbott Park
STREET: Illinois
COUNTRY: USA
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CLASSIFICATION, 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/100,708
FILING DATE: July 29, 1993
ATTORNEY/AGENT INFORMATION:
NAME: WONG, WEAR KHING
REGISTRATION NUMBER: 33,561
REFERENCE/DOCKET NUMBER: 5324.US.PI
TELECOMMUNICATION INFORMATION:
TELEFAX: (708) 938-2517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Wong, Wean Khing
REGISTRATION NUMBER: 33,561
REFERNCE/DOCKET NUMBER: 5324.US.P1
RELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: SOFTPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/657,392
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COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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BEICR APPLICATION DATA:
APPLICATION NUMBER: 08/100,708
FILING DATE: July 29, 1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acid residues
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IS-08-657-392-23
Sequence 23, Application US/08657392
Patent No. 5843634
GENERAL INPORMATION:
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Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 RIQRGPGRAFVT 13
                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
TOPOLOGY: unknown
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STATE: Illinois
COUNTRY: USA
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APPLICANT:
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APPLICANT:
APPLICANT:
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                                                                    Gaps
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                                  Length 13;
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                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                    And Their
Conjugates For Diagnostic Assays
                                                                                                                                                                                                                                                                                                                                               March, S.C.
Russell, J.R.
Yue, V.T.
VENTION: Genetically Engineered Enzymes
                              80.5%; Score 62; DB 5; I
100.0%; Pred. No. 0.0017;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SUFTWARE: SOFEPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02539
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: SOFTPC
                                                                                                                                                                                                                Sequence 20, Application PC/TUS9402539 GENERAL INFORMATION: APPLICANT: Brate, E.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: ABBOTT LABORATORIES
One Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acid residues
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Wong, Wean Khing
REGISTRATION NUMBER: 33,561
REFERENCE/DOCKET NUMBER: 5324.
TELECOMMUNICATION INFORMATION:
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                             Query Match 80.5
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                  1 RIGRGPGRAFVT 12
                                                                                                                      2 RIGRGPGRAFVT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LA
                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unknown
                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (108)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
PCT-US94-02539-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM:
PCT-US94-02539-20
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RESULT 200

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Gaps
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                                                                                                                                                                                                                       TITLE OF INVENTION: Genetically Engineered Enzymes TITLE OF INVENTION: And Their TITLE OF INVENTION: Conjugates For Diagnostic Assays NUMBER OF SEQUENCES: 34 CORRESPONDENCE ADDRESSS: ADDRESSES: ABBOTT LABORATORIES STREET: One Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 62; DB 5; 1 Pred. No. 0.0017;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: SOFTPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02539
FILING DATE:
Sequence 21, Application PC/TUS9402539 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCT-US94-02539-23; Sequence 23, Application PC/TUS9402539; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Wong, Wean Khing
REGISTRATION NUMBER: 33,561
REFERENCE/DOCKET NUMBER: 5324
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 938-3517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  : (708) 938-3517
(708) 938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                  Russell, J.R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                           Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unknown
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Best Local Similarity
Matches 12; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
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APPLICANT: Russell, J.R.
APPLICANT: Yus, V.T.
TITLE OF INVENTION: Genetically Engineered Enzymes
TITLE OF INVENTION: And Their
TITLE OF INVENTION: And Their
TITLE OF INVENTION: Conjugates For Diagnostic Assays
NUMBER OF SEQUENCES: 34
CORRESPONDENCE ADDRESSE:
ADDRESSEE: ABBOTT LABORATORIES
STREET: One Abbott Park Portional Control of CITY: Abbott Park
STATE: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    80.5%; Score 62; DB 2; Length 15; 100.0%; Pred. No. 0.0019; rive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Preu. ...
                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PELING DATE: July 29, 1993
ATTORNEY/AGENT INFORMATION:
NAME: WONG, WEAR KAING
REGISTRATION NUMBER: 33,561
TELECOMMUNICATION INPORMATION:
TELEFAX: (708) 938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: SOFTPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02539
                                                                                         SOFTWARE: SOFTPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/657,392
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCT-US94-02539-24; Sequence 24, Application PC/TUS9402539; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 15 amino acid residues TYPE: amino acid
        Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIORGPGRAFVT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
           MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-657-392-24
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; Sequence 24, Application US/08657392
; Patent No. 5843634
; GENERAL INFORMATION:
    APPLICANT: Braidon, D.P.
    APPLICANT: Bridon, D.P.
    APPLICANT: Jaffe, K.D.
    APPLICANT: Mandecki, W.
    APPLICANT: Mandecki, M.
    APPLICANT: Mandeck
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         80.5%; Score 62; DB 5; Lengtu ...
                                   And Their
Conjugates For Diagnostic Assays
     Genetically Engineered Enzymes
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                                                                                                                                                                                                                                                                                      ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM C COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: SoftPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02539
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acid residues
TYPE: amino acid
STRANDEDNES:
TOPOLOGY: unknown
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
TITLE OF INVENTION: Genetically Engrittle OF INVENTION: And Their TITLE OF INVENTION: Conjugates For NUMBER OF SEQUENCES: 34 CORRESPONDENCE ADDRESS: ADDRESSER: ABBOTT LABORATORIES STREET: One Abbott Park Road CITY: Abbott Park Road CITY: Allinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Wong, Wean Khing
REGIESTRATION NUMBER: 33,561
REFRENCE/DOCKET NUMBER: 5324
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 938-3517
TELEFAX: (708) 938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 12; Conservative
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: I.
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TITLE OF INVENTION: Genetically Engineered Enzymes And Their
TITLE OF INVENTION: Conjugates For Diagnostic Assays
NUMBER OF SEQUENCES: 34
                                                                                                                                   80.5%; Score 62; DB 3; Length 20; 100.0%; Pred. No. 0.0025; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: SOFTPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/657,392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/100,708
FILING DATE: JULY 29, 1993
ATTORNEY/AGENT INFORMATION:
NAME: WOOG, WEAN Khing
REGISTRATION NUMBER: 33,561
                                                                                                                                                                                                                                                                                                                                                                           Sequence 27, Application US/08657392
Patent No. 5843634
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3: ABBOTT LABORATORIES
One Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 23 amino acid residues
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 5.
TELECOMMUNICATION INFORMATION
TELEPHONE: (708) 938-3517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (708) 938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Jaffe, K.D.
APPLICANT: Mandecki, G.A.
APPLICANT: Mandecki, W.
APPLICANT: Russell, J.R.
APPLICANT: Russell, J.R.
APPLICANT: Yue, V.T.
TITLE OF INVENTION: Geneti
                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                          1 RIQRGPGRAFVT 12
                                                                                                                                                                                                                                                                   9 RIORGPGRAFVT 20
                                               TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-973-551-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LA
STREET: One Abbott P
      TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Abbott Park
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Abbott Park
STATE: Illinois
COUNTRY: USA
ZIP: 60064-3500
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MOLECULE TYPE: 1
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                       US-08-657-392-27
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US-08-657-392-27
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APPLICANT: Le Contel, Carole
APPLICANT: Le Contel, Carole
APPLICANT: Le Contel, Carole
APPLICANT: Le Galea, Pascale
TITLE OF INVENTION: VACCINE AGAINST INFECTIOUS AGENTS HAVING
TITLE OF INVENTION: AN INTRACELLULAR PHASE, COMPOSITION FOR THE TREATMENT A
TITLE OF INVENTION: DIAGNOSIS
TITLE OF INVENTION: DIAGNOSIS
CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,551
FILING DATE: 30-DEC-1997
CLASSIETCATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/FR96/01006
FILING DATE: 28-JUN-1996
PRIOR APPLICATION NUMBER: FR 9507914
                                                                                                                                                                                                                                                                                                                                                                                           80.5%; Score 62; DB 5; Le
100.0%; Pred. No. 0.0019;
:ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 : 3000 K Street, N.W., Suite 500
Washington
NAME: Wong, Wean Khing
REGISTRATION NUMBER: 33,561
REPERBUCE/DOCKET NUMBER: 5324.PC.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 938-2523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              65691/130
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COUNTRY: USA
ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                               INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acid residues
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-973-551-24
; Sequence 24, Application US/08973551
; Parent No. 6113902
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 30-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Granados, Patricia D.
RECISTRATION NUMBER: 33,683
REFERENCE/DOCKET NUMBER: 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (202) 672-5399
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 RIORGPGRAFVT 14
                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                unknown
                                                                                                                                                                                                                                                                                                                                             PCT-US94-02539-24
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APPLICANT: Pendleton, Careen
APPLICANT: Roslowski, Steven
APPLICANT: Roslowski, Steven
APPLICANT: Marguiles, David H.
TITLE OF INVENTION: Cytotoxic T Lymphocytes Specific for the HIV-I Envelope
NUMBER OF SEQUENCES:
ADDRESSES: Birch, Stewart, Kolash & Birch
STREET: 301 N. Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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/note= "Active peptide of HIV-I envelope
from strain IIIB"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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Pred. No. 0.0024;
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US-080-6
US-080-11-080-6
Sequence 6, Application 08/111080
Patent No. 5558865
GENERAL INFORMATION:
APPLICANT: Ohno, Teuneya
TITLE OF INVENTION: HIV Immunotherapeutics
WUMBER.OF SEQUENCES: 38
CORRESPONDENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Borun
ADDRESSEE: Borun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUBBER:
FILING DATE: 06-MAR-1992
CLASSIFICATION 435
ATTONERY/AGENT INFORMATION:
NAME: SVENESON, Leonard R.
REGISTRATION NUMBER: 30,330
REFERENCE/DOCKET NUMBER: 1173-392P
TELEPRANTION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPRAN: 703-241-1300
INPORMATION FOR SEQ ID NO: 20:
SEQUENCE CHRACTERISTICS:
LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FRAGMENT TYPE: internal ORIGINAL SOURCE: ORGANISM: Human Immunodeficiency Virus Type I STRAIN: IIIB
                                                                                                                                                  ADDRESSEE: Birch, Stewart, Kolash & Birch STREET: 301 N. Washington CITY: Falls Church STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                79.2%; Sco...
100.0%; Pred. No. ...
0; Mismatches
                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                               ZIP: 22046-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 IORGPGRAFVTI 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Peptide
LOCATION: 1.13
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDALL
STREET: 650C
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-07-847-311A-20
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                                                                                                                                                                                        APPLICANT: VaILE, K.D.

APPLICANT: Krafft, G.A.

APPLICANT: Mandecki, W.

APPLICANT: March, S.C.

APPLICANT: Yue, V.T.

ITLE OF INVENTION: Genetically Engineered Enzymes

ITLE OF INVENTION: Conjugates For Diagnostic Assays

NUMBER OF SEQUENCES: 34

CORRESPONDENCE ADDRESS:

ADDRESSEE: ABBOTT INFOL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 62; DB 5; Length 23; Pred. No. 0.0029; 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: WOIG, WEAN Khing
REGISTRATION NUMBER: 33,561
REFERENCE/DOCKET NUMBER: 5324.PC.01
TELERPONNINICATION INFORMATION:
TELERPONE: (708) 938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: SoftPC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02539
FILING DATE:
                                                                                                             PCT-US94-02539-27; Sequence 27, Application PC/TUS9402539; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NSCOL 197-311A-20
Sequence 20, Application US/07847311A
Patent No. 5976541
GENERAL INFORMATION:
APPLICANT: Berzofeky, Jay A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: ABBOTT LABORATORIES
STREET: One Abbott Park Road
CITY: Abbott Park
STATE: 1111inois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           80.55,
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: unknown
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 12; Conservative
1 RIORGPGRAFVT 12
                    1 RIQRGPGRAFVT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12 RIORGPGRAFVT 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Bicknell
STREET: Two First National Plaza, 20 South Clark
STREET: Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/07111
FILING DATE: 19920824
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                              0.0037;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 6, Application PC/TUS9207111
GENERAL INFORMATION:
APPLICANT: Ohno, Tsuneya
TITLE OF INFORTION: HIV Immunotherapeutics
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Mismatches
      US 08/039,457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/748,562
FILING DATE: 22-AUG-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 60603
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
PELICATION NUMBER: US 08/039, PELICATION NUMBER: US 08/039, PILOS DATE: 22-APR-1993, ATTORNEY/AGENT INFORMATION: NAMB: BOTUN, MICHAEL E. REGISTRATION NUMBER: 25,447, REPERRINCE/DOCKET NUMBER: 3162; TELECOMMUNICATION INFORMATION: TELEPHONE: (312) 474-6300 TELEFAX: (312) 474-646 TELEFAX: 25-3856 INFORMATION OR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS: LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 31(
THE ECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (312) 346-5750
TELEFAX: (312) 984-9740
TELEX: 25-385
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 IQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 IRIGPGRAFVTIGK 14
                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 85.7%
...rnhes 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Noland, Greta E
                                                                                                                                                                                                                                                                                                                     , MOLECULE TYPE: peptide US-08-211-980-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , MOLECULE TYPE: peptide PCT-US92-07111-6
                                                                                                                                                                                                                                                                     TOPOLOGY: 14-
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STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: AMINO ACID
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Sequence 6, Application US/08211980

Patent No. S6655569

GENERAL INFORMATION:

APPLICANT: Ohno, Tsuneya

TITLE OF INVENTION: HIV Immunotherapeutics

NUMBER OF SEQUENCES: 38

CORRESPONDENCE ADDRESS:

ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Borun
                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 08/111,080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/211,980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 6300 Sears Tower, 233 S. Wacker Drive CITY: Chicago STATE: Illinois COUNTRY: USA
                                                                                                                                                          PILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/748,562
FILING DATE: 22-NuG-1991
PRIOR APPLICATION NUMBER: PCT/US92/07111
FILING DATE: 24-NuG-1992
PRIOR APPLICATION NUMBER: US 08/039,457
FILING DATE: 24-ANG-1993
ATTORNEY/AGENT INFORMATION:
NAME: BOTUN, Michael F.
REGISTRATION NUMBER: 25,447
REFERENCE/DOCKET NUMBER: 25,447
REFERENCE/DOCKET NUMBER: 31629
TELECOMMUNICATION INFORMATION:
THE ECOMMUNICATION INFORMATION:
THE ECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIPICATION: 424
PRIOR APPLICATION DATE:
FILING DATE: 24-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               77.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 IORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 85.7
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 14 amino acids
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-111-080-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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us-08-869-386-1.rai

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US-08-704-170-74

| Sequence 74, Application US/08704170
| Sequence 74, Application US/08704170
| Patent No. 5707626
| GENERAL INFORMATION:
| APPLICANT: Takehana, Yoshi
| APPLICANT: Ehremann, Yoshi
| APPLICANT: Ehremann, THERAPEUTIC STRATEGIES FOR
| TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
| NUMBER OF SEQUENCES: 121
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Robbins, Berliner & Carson
| STREET: 201 No. 5707626th Figueroa Street, Suite 500
| CITY: Los Angeles
| STREET: California
| COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                75.3%; Score 58; DB 1; Length 11; 100.0%; Pred. No. 0.006;
                     NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
SADDRESSER: Robbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite 500
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PLOPS HS.
COMPUTER: PLOPS HS.
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,170
FILING DATE:
CLASSIPLCATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR.1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OFFICE OFFI
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COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/104,170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
RAPPLICATION NUMBER: US 08/029,850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 RGPGRAFVTIG 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 11; Conserva
                                                                                                                                                                                                                                     90012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGIH:
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Patent No. 5707626
GENERAL INFORMATION:
APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
APPLICANT: Takehana, Glenn
TITLE OF INVENTION: THEREPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
85.7%; Pred. No. 0.0037; wiematches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 5; Length 14;
                                                                                                                                                                                                                                                                                                                   Sequence 6, Application PC/TUS9307967
GENERAL INFORMATION:
APPLICANT: Ohno, Teuneya
TITLE OF INVENTION: HIV Immunotherapeutics
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marehall, O'Toole, Gerstein, Murray & STREET: 6300 Sears Tower, 233 S. Wacker Drive
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/07967
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Mismatches
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85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAS. (312) 474-0448
TELEFAS. 25-3856
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (312) 474-6300
                                                                                                                                                     2 IQRGPGRAFVTIGK 15
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                           Best Local Similarity 85.7
Matches 12; Conservative
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Best Local Similarity 85.7
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                 RESULT 211
PCT-US93-07967-6
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APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
APPLICANT: Takehana, Yoshi
TITLE OF INVENTION: THERAPBUTIC STRATEGIES FOR
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 75.3%; Score 58; DB 5; Length 11; 100.0%; Pred. No. 0.006; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/08090148
Patent No. 5534257
GENERAL INFORMATION:
APPLICANT:
APPLICANT: Stockley, Peter George
APPLICANT: Talbot, Simon John
TITLE OF INVENTION: Antigen-Presenting Capsid with
TITLE OF INVENTION: Fusion MS2-Coat Protein
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02631
                                                                                                                                                                                                                                                                                                                                               SSE: Robbins, Berliner & Carson
F: 201 North Figueroa Street, Suite 500
Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Preu....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNDER:
APPLICATION UNDER:
ILING DATE:
ILING DATE:
NAME:
Spieals, John P.
REGISTRATION NUMBER:
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
                                                                                                                                                      Sequence 74, Application PC/TUS9402631
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Rosenman & Colin
575 Madison Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213) 977-1001
TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 74
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100
Matches 11; Conservative
                                               1 RGPGRAFVTIG 11
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                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angele
STATE: Californi
COUNTRY: U.S.A.
ZIP: 90012
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GY: linear
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STREET: 575
CITY: New Y
                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
STREET: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCT-US94-02631-74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 216
US-08-090-148-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
                                                                                                                                                                                                                                                                                                                                                      Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.3%; Score 58; DB 5; Length 11; 100.0%; Pred. No. 0.006; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: STATEM PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
RAPPLICATION NUMBER: PCT/US94/02631
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS: 121
ADDRESSEB: Robbins, Berliner & Carson
STREET: 201 North Figueroa Street, Suite 500
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                    DB 1;
0.006;
                                                                                                                                                                                                                                                                                                                                               Query Match 75.3%; Score 58; DB Best Local Similarity 100.0%; Pred. No. 0.0 Matches 11; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DALL.
CLASIPICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
FREFERNEK/ECOKCET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
""" REPERENCE/DOCKET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
""" REPERENCE/DOCKET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
""" ATT-1003
                                                                                      1920-331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 73, Application PC/TUS9402631
; GENERAL INFORMATION:
             ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFRENCE/DOCKET NUMBER: 1920-
TELECOMMUNICATION INFORMATION:
TELEFAX: (213) 977-1001
TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 74:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
FILING DATE: 11-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                               ; MOLECULE TYPE: peptide US-08-704-170-74
                                                                                                                                                                                                                                                                                                                                                                                                                                         4 RGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RGPGRAFVTIG 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide PCT-US94-02631-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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PCT-US94-02631-73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Virginia
COUNTRY: USA
CONDUTER: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,277
FILING DATE: 06-Jun-1995
CLASSIFICATION: cUnknown>
                                                                                                                                                                                                                       Score 58; DB 1; Length 15;
Pred. No. 0.0081;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 58; DB 3; Length 15;
Pred. No. 0.0081;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10, Application US/08466277

Patent No. 6190666

GENERAL INFORMATION:

APPLICANT: Garoff, Henrik

Liljestrom, Peter

Illjestrom, Repression Systems Based on Alphaviruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/920,281
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: MUTPHY Jr., Gerald M.
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 828-103P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 703-241-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 703-241-2848
                                               10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 75.3%;
Best Local Similarity 84.6%;
Matches 11; Conservative
                                                                                                                                                                                                                            Query Match
Best Local Similarity 84.6%;
Matches 11; Conservative
              TELEX: 248345
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                         1 RIORGPGRAFVTI 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 RIQRGPGRAFVTI 13
                                                                                                                                                                                                                                                                                                                                                                 3 RIORGPGRAFVEL 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 RIORGPGRAFVEL 15
  703-241-2848
                                                                                                                             ; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-07-920-281C-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 248345
                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-466-277-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-466-277-10
TELEFAX:
TELEX: 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10. Application US/07920281C

Sequence No. 5739026

GENERAL INFORMATION:
APPLICANT: Garoft, Henrik
TITLE OF INVENTION: DNA Expression Systems Based on
TITLE OF INVENTION: Alphaviruses
TITLE OF INVENTION: Alphaviruses
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747

CITY: Falls Church
STRATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 22040-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/920,281C
FILING DATE: 13-AUG-1992
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MALPHY JT., Gerald M.
REGISTRATION NUMBER: 28,977
                          COUNTRY: U.S.A.
ZIP: 1002-2585
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5", 1.44Mb
COMPUTER: IBM PS2.466
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/090,148
FILING DATE: 08/11/93
CLLASSIPETCATION BATA:
APPLICATION NUMBER: GB 9101550.3
FILING DATE: 01/24/91
APPLICATION NUMBER: PCT/GB92/00124
FILING DATE: 01/24/91
APPLICATION NUMBER: 27,582
ATTORNEY/AGENT INFORMATION:
NAME: Nissenbaum, Israel
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (212) 940-6404
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           828-103P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 8:
TELECOMMUNICATION INFORMATION
TELEPHONE: 703-241-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 13 AMINO ACIDS
TYPE: AMINO ACID
TOPOLOGY: NOT RELEVANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 GPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , MOLECULE TYPE: PEPTIDE US-08-090-148-5
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us-08-869-386-1.rai

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Sequence 70, Application US/08704170

| Sequence 70, Application US/08704170
| Patent No. 5707626
| GENERAL INFORMATION:
| APPLICANT: Davas, Angeline APPLICANT: Takehana, Yoshi
| APPLICANT: Ehreaman, Glenn
| APPLICANT: Ehreaman, Glenn
| TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
| NUMBER OF SEQUENCES: 121
| CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSEE: Robbins, Berliner & Carson
| STREET: 201 No. 5707626th Figueroa Street, Suite 500
| CITY: Los Angeles
| STATE: California | COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    74.0%; Score 57; DB 5; Length 11; 100.0%; Pred. No. 0.0086;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,170
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: 1BM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: 1BM P.C. DOS (Version 3.30)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/06688
FILING DATE: 19920811
                                                                                                                          PELICALIUM DATE: 19920811

FILING DATE: 19920811

CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 744,281

FILING DATE: 13 August 1992
ATTORNEY/AGENT INFORMATION:
NAME: Paul T. Clark
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00231/052WO1
TELECOMMUNICATION INFORMATION:
TELEPPONE: (617) 542-8906
TELEPPONE: (617) 542-8906
TELEPPIONE: (617) 542-8906
TELEPPIONE: (617) 542-8906
TELEPPIONE: SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIQRGPGRAFV 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: AMINO ACID
; TOPOLOGY: linear
PCT-US92-06688-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 11; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90012
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Sequence 14, Application PC/TUS9206688

GENERAL INFORMATION:
APPLICANT: REBLIGEN CORPORATION
APPLICANT: THE ROCKEFELLER UNIVERSITY
TITLE OF INVENTION: WALTIPLE ANTIGEN PEPTIDES FOR USE AS HIV
TITLE OF INVENTION: VACCINES
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                             STATE: Virginia
COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: TBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/688,842
FILING DATA:
FRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             75.3%; Score 58; DB 4; Length 15; 84.6%; Pred. No. 0.0081;
                                                                          Sequence 10, ....
Patent No. 6770283
GENERAL INFORMATION:
APPLICANT: Garoff, Henrik
Liljestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based on Alphaviruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION UNMBER: 08/466,277
FILING DATE: «Unknown»
ATTORNEY, AGENT INFORMATION:
NAME: Murphy Jr., Gerald M.
REGISTRATION NUMBER: 28,977
TELEPHONE: 703-241-1300
TELEPHONE: 703-241-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 10:
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Best Local Similarity 84.6
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ELEX: 248345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 02110-2804
COMPUTER READABLE FORM:
                                                      -09-688-842-10
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GENERAL INFORMATION
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Sequence 70, Application PC/TUS9402631
GENERAL INFORMATION:
APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
                                                                                                                                             Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 15;
                                                                                                                                                                                 0; Indels
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0,000
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 North Figueroa Street, Suite 500
                                                                                                                             Score 57; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          74.0%; Score 57; DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Pred. No. 0.0 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 12, Application US/07920281C; Patent No. 5739026
                                                                                                                                           Query Match

Best Local Similarity 100.0%; P:
Matches 11; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) 977-1001
TELEPAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
TYPE: amino acids TOPOLOGY: 14-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 11; Conservative
            SEQUENCE CHARACTERISTICS
                                                                                  ; MOLECULE TYPE: peptide US-08-704-170-70
                                                                                                                                                                                                                        1 RIORGPGRAFV 11
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CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-07-920-281C-12
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Gaps
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APPLICANT: Garoff, Henrik
APPLICANT: Liljestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based on
TITLE OF INVENTION: Alphaviruses
NUMBER OF SEQUENCES: 27
CORRESPONDENCE: ADDRESS:
ADDRESSE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 57; DB 1; Length 17;
Pred. No. 0.013;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Garoff, Henrik
Liljestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based on
Alphaviruses
                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

CCMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURENT APPLICATION DATA:

APPLICATION NUMBER: US/07/920,281C

FILING DATE: 13-AUG-1992

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Murphy Jr., Gerald M.

REGISTRATION NUMBER: 28,977

REFERENCE/POCKET NUMBER: 28,977

REFERENCE/POCKET NUMBER: 28,977

RELECOMMUTICATION INFORMATION:

TELECOMMUTICATION INFORMATION:

TELECOMMUTICATION INFORMATION:

TELEFRA: 703-241-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 74.0%; Scorio 100.0%; Pred. No. v. 100.0%; Pred. No. v. v. Mismatches
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APPLICATION NUMBER: US/08/466,277
FILING DATE: 06-Jun-1995
CLASSIPICATION: «UNKNOWN»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: Virginia
COUNTRY: USA
ZIP: 22040-077
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 12, Application US/08466277
Patent No. 6190666
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 703-241-2848
TELEX: 248345
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTEREGICS:
LENGTH: 17 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIQRGPGRAFV 11
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Gaps

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                                                    74.0%; Score 57; DB 4; Length 17; 100.0%; Pred. No. 0.013;
                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
                                                                                                                                                                                                                                                          US-08-257-528B-16

Sequence 16, Application US/08257528B

Patent No. 5639854

GENERAL INFORMATION:
APPLICANT: STA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Toronto
COUTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: ParentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/257,528B
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 16, Application US/08460602A
Patent No. 5759769
GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 57; DB 1;
Pred. No. 0.016;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I
REGISTRATION UNDHERE: 24,971
REFERENCE/DOCKET UNDHER: 1038
TELECOMMUNICATION INFORMATION:
TELEFONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                    Best Local Similarity 100.
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 21 amino acids
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Best Local Similarity 91.7
Matches 11, Conservative
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                                                                                                                                     1 RIQRGPGRAFV 11
                                                                                                                                                                           4 RIORGPGRAFV 14
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: linear
US-08-257-528B-16
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US-08-460-602A-16
          US-09-688-842-12
                                                  Query Match
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ADDRESSE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
STREET: P.O. Box 747
STREET: Virginia
COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATE:
PILING DATE: 17-Oct-2000
CLASSIFICATION: AUNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                        74.0%; Score 57; DB 3; Length 17; 100.0%; Pred. No. 0.013; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 12, Application US/0968842
; Patent No. 6770283
; GENERAL INFORMATION:
HAPLICANT: Garoff, Henrik
Illestrom, Peter
TITLE OF INVENTION:
Alphaviruees
               APPLICATION NUMBER: 07/920,281
FILING DATE: «Unknown»
ATTORNEY,AGENT INFORMATION:
NAME: Murphy Jr., Gerald M.
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 828-103P
TELECOMUNICATION INFORMATION:
TELEFRAX: 703-241-1300
TELEFRX: 703-241-2848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/466,277
FILING DATE: «Unknom»
ATTORNEY, AGENT INFORMATION:
NAME: Murphy Jr., Gerald M.
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 828-103P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-241-1300
TELEFAK: 703-241-1300
                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: linear; MOLECULE TYPE: protein; SEQUENCE DESCRIPTION: SEQ ID NO: 12: US-98-466-277-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                         TELEX: 24845
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 RIORGPGRAFV 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ELEX: 248345
                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
Matches 11; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 225
US-09-688-842-12
                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Gaps

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Score 57; DB 1; Length 21;
Pred. No. 0.016;
0; Mismatches 1; Indels
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18-08-465-217A-16

18-08-465-217A-16

18-08-465-217A-16

18-08-465-217A-16

18-08-21-217A-16

18-08-217A-16

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MEDIUM TYPE: Floppy disk
COMPUTER: TEM PC COMPATIBLE
COPERATIOS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN DATA:
APPLICATION NUMBER: US/08/465,217A
FILING DATE: 09-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
ATCORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-486 MIS:jb
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-UN-1994
CLASSIPTCATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-UN-1993
CLASSIFTCATION: 424
ATTONNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 1038-487 MIS:jb
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPANE: (416) 595-1163
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH. 21 main acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 91.7
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7 RIORGPGRAFYT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-463-966A-16
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Sequence 16, Application US/08463966A

Sequence 16, Application US/08463966A

PAPLICANT: S195955

APPLICANT: GLANG, Pele

APPLICANT: CHONG, Pele

APPLICANT: CHONG, Pele

APPLICANT: TILE OF INVENTION: Tandem Synthetic HIV-1 Peptides

WUMBER OF SEQUENCES: 101

CORRESPONDENCE AMCBURDA

STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSE: Suite 701, 330 University Avenue
CITY: Toronto
STATE: Onterio
CITY: Toronto
STATE: Onterio
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PEATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/463,966A
FILING DATE: 05-UN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,602A
FILING DATE: 02-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424

RELING DATE: 09-JUN-1993
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATCORNEY AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-450 MIS:jb
TELECOMMUNICATION INPORMATION:
TELEPHONE: (416) 595-1153
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 emino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 57; DB 1;
Pred. No. 0.016;
0; Mismatches
           Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 91.7%;
Matches 11; Conservative
                                                    CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7 RIORGPGRAFYT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: linear
US-08-460-602A-16
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0; Gaps

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ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Best Local Similarity 91.7
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RIQRGPGRAFVT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-462-507A-16
                                                                                                              JS-08-462-507A-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-467-881A-16
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                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                      ;
0
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                                                                                                                                                                                                                                                         74.0%; Score 57; DB 1; Length 21; 91.7%; Pred. No. 0.016;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                   1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: CLBIN, Michel H.
ITILE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Tocatio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: CALLAGAGE
ZIP: MGG LRY
ZIP: MGG LRY
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN BATA:
APPLICATION NDATA:
APPLICATION NDATA:
APPLICATION NOATA:
APPLICATION NOATA:
APPLICATION NOATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NOATA:
ATONATION FOR SECUENCE CHARACTERISTICS:
APPLICATION ACTION TO SECUENCE CHARACTERISTICS:
APPLICATION ACTION ACT
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Pred. No. 0.016;
0; Mismatches
                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 230
US-08-464-329A-16
Equence 16, Application US/08464329A
; Patent No. 5817754
.... 21 amino acide
...eE: amino acide
STRANDEDNESS: single
;
TOPOLOGY: linear
US-08-465-217A-16
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91.7%;
                                                                                                                                                                                                                                                                                                                      11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-464-329A-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 11; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ontario
7: Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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Sequence 16, Application US/08462507A

Patent No. 5876731

GENERAL INFORMATION:

APPLICANT: CHONG, Pele

APPLICANT: KLEIN, Michel H.

TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides

NUMBER OF SEQUENCES: 101

CORRESPONDENCE ADDRESS:

ADDRESSEE: Sim & McBurney

STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATION STSTEM: TC-LUCY MS-LUCS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/462,507A
FILING DATE: 05-UN-1995
CLASSIFICATION 3.424
PRIOR APPLICATION NUMBER: 08/257,528
FILING DATE: 09-UN-1994
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-UN-1994
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-UN-1993
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-UN-1993
CLASSIFICATION NUMBER: 09-440-1993
FILING DATE: 09-10-1993
CLASSIFICATION NUMBER: 1038-451 MIS:jb
FILING DATE: 09-10-1993
CLASSIFICATION NUMBER: 1038-451 MIS:jb
FILING DATE: 08-0-10-1993
CLASSIF
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Patent No. 5951986
GENERAL INFORMTON:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
```

```
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitale, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 71:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acide
TYPE: Amino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
BILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920-331
TELEPHONE: (213) 977-1001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 71, Application PC/TUS9402631 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.1
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
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MOLECULE TYPE: peptide
PCT-US94-02631-71
                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 GPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GPGRAFVTIG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 90012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCT-US94-02631-71
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GENERAL INFORMATION:

APPLICANT: Douves, Angeline

APPLICANT: Takehana, Yoshi

APPLICANT: Takehana, Yoshi

APPLICANT: Bhresmann, Glan

TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS

INUMBER OF SEQUENCES: 121

CORRESPONDENCE ADDRESS:

ADDRESSEE: Robbins, Berliner & Carson

STREET: 201 No. 5707626th Figueroa Street, Suite 500

CITY: Los Angeles

STREET: California

COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         74.0%; Score 57; DB 2; Length 21; 91.7%; Pred. No. 0.016;
                      COMPUTER READABLE FORM:

ABGINATION OF STATES COMPATION:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RC-BAS MS-DOS
SOFTWARE: PATENTIN RC-BAS MS-DOS
SOFTWARE: PATENTIN RC-BAS MS-DOS
SOFTWARE: PATENTIN RC-BAS MS-S
CLASSIFICATION NUMBER: US/08/467,881A
FILING DATE: 06-JUN-1995
CLASSIFICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: A24
PRIOR APPLICATION: A24
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 1038-488 MIS:jb
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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MEDIUM TYRER READABLE FORM:
MEDIUM TYRE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/104,170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 71, Application US/08704170 Patent No. 5707626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 21 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIQRGPGRAFVT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7 RIORGPGRAFYT 18
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         Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-467-881A-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 233
US-08-704-170-71
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Matches
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Takehana, Yoshi
APPLICANT: Takehana, Yoshi
APPLICANT: Takehana, Yoshi
APPLICANT: Takehana, Yoshi
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 North Figueroa Street, Suite 500
68.8%; Score 53; DB 1; Length 10;
100.0%; Pred. No. 0.033;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                68.8%; Score 53; DB 5; Length 10;
100.0%; Pred. No. 0.033;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READBALE FORM:
MEDIUM TYPE: Floppy disk
COMFUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02631
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-UG-25-366A-36

| Sequence 36, Application US/08463966A
| Patent No. 5795955
| GENERAL INFORMATION:
| APPLICANT: CHONG, Pele
| APPLICANT: CHONG, Pele
| APPLICANT: KIEIN, Michel H.
| TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
| NUMBER OF SEQUENCES: 101
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Sim & McBurney
| STREET: Suite 701, 330 University Avenue
| CITY: Toronto | STATE: Oncario
| COUNTRY: Canada | CITY: Toronto | STATE: Oncario | |
| CONVIRY: Canada | CONVIRY: Canada | CONVIRY: Canada | CONVIRTER: IBM PC compatible |
| COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk | COMPUTER: IBM PC compatible |
| COMPUTER: IBM PC compatible | OPERATING SYSTEM: PC-DOS/MS-DOS |
| COMPANE: Patentin Release #1.0, Version #1.25 |
| CURRENT APPLICATION DATA: |
| APPLICATION NUMBER: US/08/463,966A |
| FILING DATE: NAME | NAME |
| CLASSIFICATION DATA: |
| DATOR ADDITOTION DATA: |
| D
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOSTANTARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,602A
FILING DATE: 02-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
FILING DATE: 09-JUN-1994
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION OF 424
ATTOKNEY/AGENT INFORMATION:
NAME: STEWARRY, MICHAEL I.
REGISTRATION NUMBER: 24,973
RELEPRORE/POCKET UNBER: 24,973
RELEPRORE/POCKET UNBER: 24,973
RELEPRORE (416) 595-1155
TELEFRANCE/POCKET UNBER: 24,973
RELEPRORE (416) 595-1165
TELEFRANCE (416) 595-1165
TELEFRANCE CHARACTERISTICS:
TENDORMATION POR SEO ID NO: 36:
SEQUENCE CHARACTERISTICS:
TENDORMATION POR SEO ID NO: 36:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY:
US-08-460-602A-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RIQRGPGRAF 10
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Trad No. 0.046;
0; Indels
                                                                                                                                                                                                                                             Sequence 36, Application US/08257528B

Patent No. 5639854

GENERAL INFORMATION:
APPLICANT: CHONG, Pele
APPLICANT: CHONG, Pele
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS: 103
CITY: Toronto
STATE: Ontario
STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 09-JUN-1994
FILING DATE: 09-JUN-1994
FILING DATE: 09-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 36, Application US/08460602A

Patent No. 5759769

GENERAL INFORMATION:
APPLICANT: SIA, CHARLES D.Y.
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
CITY: Toronto
CITY: Ontario
CONTYRY: Canada

ZIP: M5G 1R7
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-336 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.08;
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INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 14 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 68.8
Best Local Similarity 100.
Matches 10; Conservative
                                    5 GPGRAFVTIG 14
                                                                                   GPGRAFVTIG 10
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                                                                                                                                                                                              RESULT 235 - US-08-257-528B-36
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Query Match
Best Local Similarity
Matches 10; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 240
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Sequence 36, Application US/08465217A

Sequence 36, Application US/08465217A

Patent No. 5800822

GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.

TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES:
ADDRESSER: SIM & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,217A
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA:
RILNG DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                          0.046;
                                                                                                                                                                                                                                                                                                                                                                              Query Match 68.8%; Score 53; DB 1; Best Local Similarity 100.0%; Pred. No. 0.046; Matches 10; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
                                                                                   1038-487 MIS:jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1038-486 MIS: jb
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,971
REFERENCE/DOCKET NUMBER: 1039.
TELECOMMUNICATION INFORMATION:
TELEFAX: (416) 595-1155
TELEFAX: (416) 595-1155
TELEFAX: (416) 595-1155
TELEFAX: LANGUERE SEQ ID NO: 36: SEQUENCE CHARACTER.ESTICS:
LENGTH: 14 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFRENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 36:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 RIQRGPGRAF 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 RIORGPGRAF 14
                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear
US-08-463-966A-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Canada
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-465-217A-36
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0
Length 14;
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                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                      US-08-464-329A-36
Sequence 36, Application US/08464329A
Patent No. 5817754
GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: SIA, Charles D.Y.
APPLICANT: Michel H.
TILLE OF SEQUENCES: 101
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,329A
FILING DATE: 05-UW-1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 53; DB 2;
Pred. No. 0.046;
Score 53; DB 1;
Pred. No. 0.046;
                                                                                                                                                                                                                                                                                                                                                                 E: Sim & McBurney
Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1038-449 MIS:jb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-462-507A-36; Sequence 36, Application US/08462507A; Patent No. 5876731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-UUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INPORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFRENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 10; Conservative
                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
                                                                                                         5 RIQRGPGRAF 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 RIORGPGRAF 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: linear
US-08-464-329A-36
                                                                                                                                                                                                                                                                                                                                                                     Abbar
STREET: Sur-
TTTY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: Ontario
COUNTRY: Canada
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RESULT 242
US-08-218-025A-16
i Sequence 16, Application US/08218025A
j Sequence 16, Application US/08218025A
j Sequence 16, Application US/08218025A
j Settent No. 5556744
j GENERAL INFORMATION;
j APPLICANT: Weiner, David B.
j APPLICANT: Weiner, William V.
j TITLE OF INVENTION: Methods and Compositions for Diagnosing
TITLE OF INVENTION: Methods and Compositions for Diagnosing
TITLE OF INVENTION: Methods and Compositions for Diagnosing
TITLE OF INVENTION: Methods and Compositions for Diagnosing
TITLE OF INVENTION: Methods and Treating Certain HIV Infected Patients
NUMBER OF SEQUENCES: 197
CORRESPONDENCE ADDRESS:
ADDRESSEE: Howson and Howson
STREET: P.O. Box 457, 321 No. 5556744ristown Road
CITY: Spring House
j STATE: Pennsylvania
country: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: ABC and a compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: ABC and a compatible

COMPUTER: ABC and a compatible

CLASSIFICATION NUMBER: US/08/218,025A

FILING DATE: 24-MAR.1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/891,451

FILING DATE: 39-MAY-1992

ATTORNEY/AGENT INFORMATION:
                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                            CUKKENT AFPLICATION NUMBER: US/08/467,881A
PILING DATE: 06-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION NUMBER: 24,973
REGISTRATION NUMBER: 24,973
REGISTRATION NUMBER: 24,973
REGISTRATION NUMBER: 24,973
REGISTRATION NUMBER: 1038-488 MIS:jb
TELEPHONE: (416) 595-1165
TELEPHONE: (416) 595-1165
TELEPHORE: (416) 595-1165
TELEPHORE: (416) 595-1165
TELERAX: (416) 595-1165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 53; DB 2;
Pred. No. 0.046;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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REGISTRATION NUMBER: 31,215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: WE FELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 68.8%; Best Local Similarity 100.0%; Matches 10; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-467-881A-36
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CARPLICANT: CHONG, Pele
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SUGUENCES: 101
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREES: Suit 701, 330 University Avenue
CITY: Toronto
STATE: Ontario
COUNTY: Canada
ZIP: MSG IR7
COMPUTER PENTIT
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Sequence 36, Application US/08467881A

Sequence 36, Application US/08467881A

Sequence 36, Application US/08467881A

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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/462,507A
FILING DATE: 05-JUN-1995
CLASSIFICATION DATA:
APPLICATION DATA: 06-JUN-1994
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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Best Local Similarity 100.0%; Pi
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 14 amino acids
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US-08-462-507A-36
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Gaps

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GENERAL INFORMATION:
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Pred. No. 0.064;
1; Mismatches 1; Indels
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ZUDI 02110-2804

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

COMPUTER: IBW PS/2 Model 502 or 55SX

OPERATING SYSTEM: IBM P.C. DOS (Version 3.30)

SOFTWARE: WordPerfect (Version 5.0)

CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/01303

FILING DATE: 19-20214

CLASSIFICATION DATA:
APPLICATION NUMBER: 07/655,669

FILING DATE: 14-FRB-1991

ATTORNEY/AGENT INFORMATION:
NAME: PAUL T.CLARK
REGISTRATION NUMBER: 00231/050W01

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 12, Application PC/TUS9201303
GENERAL INFORMATION:
APPLICANT: Murray, Michael G. et al.
TITLE OF INVENTION: POLIOVIRUS-BASED VACCINES
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                 Query Match 68.8%; Score 53; DB 1;
Best Local Similarity 100.0%; Pred. No. 0.049;
Matches 10; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELERX: (617) 542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 12:
                             TELEPAX: (215) 540-5818
INPORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acida
(215) 540-9206
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Best Local Similarity 83.3%;
Matches 10; Conservative
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                                                                                                                                                                                                                                    ; MOLECULE TYPE: peptide
US-08-218-025A-16
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Massachusetts
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                                                                                                                                                                                                              unknown
                                                                                                                                                                      TYPE: amino acid
TOPOLOGY: unknown
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PCT-US92-01303-12
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TELEPHONE:
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US-09-820-484-8
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JUS-09-430-470-24

j Sequence 24, Application US/09430470

j Patent No. 656200

j GENERAL INFORMATION:
    APPLICANT: WOMILIAN, Minnie
    TITLE OF INVENTION: THE USE OF IMMUNPOTENTIATING SEQUENCES
    TITLE OF INVENTION: POR INDUCING IMMUNE RESPONSE
    TITLE OF INVENTION: POR INDUCING IMMUNE RESPONSE
    TITLE OF INVENTION: POR INDUCING IMMUNE RESPONSE
    CURRENT APPLICATION NUMBER: US/09/430,470
    CURRENT FILING DATE: 1999-10-29
    BARLIER APPLICATION NUMBER: US 60/106,506
    SARLIER FILING DATE: 1999-10-30
    NUMBER OF SOI ID NOS: 25
    SOFTWARE: FRAESEQ for Windows Version 4.0
    TENDENT 1.00 24
APPLICANT: Raz, Eyal
APPLICANT: Raz, Eyal
APPLICANT: Ro, Hearn Jay
APPLICANT: Richman, Douglas
APPLICANT: Richman, Douglas
APPLICANT: Horner, Anthony A.
TITLE OF INVENTION: Lymphocyte Response in vivo.
TITLE OF INVENTION: Lymphocyte Response in vivo.
FILE REFERENCE: 06510-188019.
CURRENT APPLICATION NUMBER: US 60/192,537
FRICR FILING DATE: 2001-03-28
PRIOR FILING DATE: 2000-03-28
PRIOR FILING DATE: 2000-03-28
PRIOR FILING DATE: 2000-05-11
PRIOR PILING DATE: 2000-05-11
PRIOR FILING DATE: 2000-05-11
PRIOR FILING DATE: 2000-05-11
PRIOR FILING DATE: 2000-05-11
SRIOR FILING DATE: 2000-05-11
SRIOR FILING DATE: 2000-05-11
FRIOR FILING DATE: 2000-05-11
SRIOR FILING DATE: 2000-05-11
FRIOR FILING DATE: 2000-05-11
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100.0%; Pred. No. 0.047;
tive 0; Mismatches 0; IndelB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , OTHER INFORMATION: HIV-1 class I-restricted gp120 peptide
US-09-820-484-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Human immunodeficiency virus (HIV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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; Patent No. 6592872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 10; Conservative
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APPLICANT: MCMichael, Andrew
APPLICANT: Hill, Adrian V.S.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Gilbert, Jorg
APPLICANT: Schneider, Jorg
APPLICANT: Schneider, Jorg
APPLICANT: Plebanski, Madalena
APPLICANT: Hanke, Tomas
APPLICANT: Hanke, Tomas
APPLICANT: Blanchard, Tom
APPLICANT: Blanchard, Tom
APPLICANT: Mith Geoffrey L.
APPLICANT: Mith Generate A CD8 T Cell Immune Response
TITLE OF INVENTION: Which Generate A CD8 T Cell Immune Response
FILE REFERENCE: 2907.100-000
CURRENT APPLICATION NUMBER: US/09/454,204A
FRIOR APPLICATION NUMBER: ETJGB98/01681
PRIOR APPLICATION NUMBER: GB 97 11957.2
PRIOR FILING DATE: 1997-06-09
NUMBER OF SEQ ID NOS: 78
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 68
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Pacent No. 6749856
APPLICANT: Berzofsky, Jay A. APPLICANT: Belyakov, Igor M. APPLICANT: Derby, Michael A. APPLICANT: Kelsall, Brian L. APPLICANT: Strober, Warren APPLICANT: THE GOVERNMENT OF THE UNITED STATES OF AMERICA, as
                                                                                                                                                                                                                                                                                                                         67.5%; Score 52; DB 4; Length 10;
100.0%; Pred. No. 0.047;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   67.5%; Score 52; DB 4; Length 10;
100.0%; Pred. No. 0.047;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: CTL Peptide Epitope of HIV gag
US-09-454-204A-68
                                                                                                                                                                                                                                               ; OTHER INFORMATION: CTL Epitope of HIV-1 gp120 US-09-454-204A-51
                              PRIOR FILING DATE: 1997-06-09
NUMBER OF SEQ ID NOS: 78
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 10
           PRIOR APPLICATION NUMBER: GB 97 11957.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 68, Application US/09454204A
Patent No. 6663871
                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 10; Conservative
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Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                     4 RGPGRAFVTI 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                  TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-454-204A-68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-508-552-16
                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 249
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                                                                                          Berzofsky, Jay A.
Berzofsky, Jay A.
TITLE OF INVENTION: Targeting Antigens to the WHC Class I
Processing Pathway With an Anthrax Toxin Fusion Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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| Patent No. 6663871
| GENERAL INFORMATION:
| APPLICANT: McMichael, Andrew
| APPLICANT: McMichael, Andrew
| APPLICANT: McMichael, Andrew
| APPLICANT: Hanke, Tomas
| APPLICANT: Plebanski, Magdalena
| APPLICANT: Smith, Geoffrey L.
| APPLICANT: Smith, Geoffrey L.
| APPLICANT: Mich, Geoffrey L.
| TITLE OF INVENTION: Which Generate A CD8 T Cell Immune Response
| TITLE OF INVENTION: Which Generate B CD8 T Cell Immune Response
| TITLE OF INVENTION: Which Generate B CD8 T Cell Immune Response
| TURE REPERENCE: 2997.100-000
| CURRENT APPLICATION NUMBER: US/09/454,204A
| CURRENT FILING DATE: 1998-12-09
| PRIOR PPLICATION NUMBER: PCT/GB98/01681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
APPLICATION NUMBER: US/08/937,276A
FILING DATE: 15-Sep-1997
CLASSIFICATION SUMBER: US 60/025,270
FILING DATE: INTERPLIGHT
APPLICATION NUMBER: US 60/025,270
FILING DATE: INTERPLIGHT
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 67.5%; Score 52; DB 4; Length 10; Best Local Similarity 100.0%; Pred. No. 0.047; Matches 10; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Weber, Kenneth A.
REGISTRATION NUMBER: 31,677
REFRENCE/DOCKET NUMBER: 015280-290100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                              Goletz, Theresa J.
Arora, Naveen
                                                                                  Leppla, Stephen H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 10 amino acids
APPLICANT: Klimpel, Kurt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RGPGRAFVTI 10
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FILE REFERENCE: 368200PCSEQ;
CURRENT APPLICATION: MUCOSAL CYTOTOXIC T LYMPHOCYTE RESPONSES;
CURRENT APPLICATION NUMBER: US/09/508,552;
CURRENT FILING DATE: 2000-06-12;
PRIOR PILING DATE: 1997-09-11;
PRIOR PILING DATE: 1997-09-11;
PRIOR PILING DATE: 1997-09-11;
PRIOR PILING DATE: 1998-02-17;
NUMBER OF FELING DATE: 198-02-17;
NUMBER OF FEU ID NOS: 2.0;
SOFTWARE: PATENTIN Ver. 2.0;
SEQ ID NO 16;
I.ENGTH.
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67.5%; Score 52; DB 4; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.047;
Matches 10; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application PC/TUS9201303
GENERAL INFORMATION:
APPLICANT: Murray, Michael G. et al.
APPLICANT: Murray, Michael G. et al.
TITLE OF INVENTION: POLIOVIRUS-BASED VACCINES
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORPATE: Massachusetts
COUNTRY: U.S.A.
COUNTRY: U.S.A.
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5° Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 502 or 555X
COMPUTER: IBM PS/2 Model 502 or 555X
COMPUTER: WordPerfect (Version 3.30)
SOFTWARE: WordPerfect (Version 5.0)
CURRENT APPLICATION NUMBER: PCT/US92/01303
FILING DATE: 19920214
CLASSIFICATION: c.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 10
TYPE: PRT
ORGANISM: Human immunodeficiency virus type 1
US-09-508-552-16
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CLASSIFICATION: 564
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/655,669
FILING DATE: 14-FEB-1991
ATTORNEY/AGENT INFORMATION:
NAME: PAUL T.CLARK
REGISTRATION WUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00231/05
TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUNICATION: TELECOMMUN
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'INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (617) 542-5070 TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 RGPGRAFVTI 13
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; TOPOLOGY: linear
PCT-US92-01303-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        셤
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Search completed: May 16, 2005, 14:41:20 Job time : 45 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

May 16, 2005, 12:57:08 ; Search time 70.7692 Seconds (without alignments) 70.804 Million cell updates/sec

US-08-869-386-1 Title: Perfect score:

1 RIQRGPGRAFVTIGK Scoring table: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5

1432185 segs, 334051727 residues Searched:

Total number of hits satisfying chosen parameters:

325800

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

•• Database

Published Applications AA:\*

1: /cgn2 6/prodata/1/pubpaa/USO7 PUBCOMB.pep:\*
2: /cgn2 6/prodata/1/pubpaa/PCT NEW PUB.pep:\*
3: /cgn2 6/prodata/1/pubpaa/USO6 NEW PUB.pep:\*
4: /cgn2 6/prodata/1/pubpaa/USO6 NEW PUB.pep:\*
5: /cgn2 6/prodata/1/pubpaa/USO6 NEW PUB.pep:\*
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7: /cgn2 6/prodata/1/pubpaa/USO8 NEW PUB.pep:\*
8: /cgn2 6/prodata/1/pubpaa/USO8 NEW PUB.pep:\*
9: /cgn2 6/prodata/1/pubpaa/USO8 NEW PUB.pep:\*
10: /cgn2 6/prodata/1/pubpaa/USO8 PUBCOMB.pep:\*
11: /cgn2 6/prodata/1/pubpaa/USO9 NEW PUB.pep:\*
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16: /cgn2 6/prodata/1/pubpaa/USO0 PUBCOMB.pep:\*
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18: /cgn2 6/prodata/1/pubpaa/USIO PUBCOMB.pep:\*
19: /cgn2 6/prodata/1/pubpaa/USIO NEW PUB.pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	Sequence 15, Appl	Sequence 24, Appl	Sequence 8, Appli	Sequence 9, Appli	Sequence 3, Appli	Sequence 246, App	Sequence 262, App	Sequence 6, Appli	Sequence 2, Appli	Sequence 44, Appl	Sequence 3, Appli	Sequence 3, Appli	Sequence 25, Appl
QI 1	US-09-810-310-15	US-09-810-310-24	US-09-989-621-8	.0 US-09-827-688-9	.0 US-09-077-439A-3	4 US-10-133-210-246	4 US-10-133-210-262	4 US-10-147-910-6	7 US-10-787-880-2	4 US-10-062-710-44	ני	15 US-10-283-610A-3	14 US-10-178-488-25
* Query Match Length DB ID	15 9	15 9	15 9	15 1	15 1	15 1	15 1	15 1	15 1	16 1	20 9	20	21
Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
Score	77	77	77	77	77	77	77	77	77	77	77	77	77
Regult No.	-	~	e	4	S	ø	7	σ.	O	10	11	12	13

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101
14 US-10
14 US-1(
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10 US-09
14 US-10
16 US-10-398
9 US-09-901
14 US-10-2
17 US-10-622
9 US-09-901
-60-SD 6
9 US-09-810
-60-SD 6
9 US-09-087
9 US-09-087-513-1
10 US-09-
10 US-09
14 US-10
14 US-1
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14 US-1
14 US-1
14 US-1
14 US-1
14 US-10

## ALIGNMENTS

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Gaps
                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: HIV PEPTIDE; OTHER INFORMATION: ANTIGEN
US-09-810-310-15
                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                      Query Match
100.0%; Score 77; DB 9; Length 15;
Best Local Similarity 100.0%; Pred. No. 1.9e-05;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
US-09-810-310-15
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1 RIORGPGRAFVTIGK 15

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RESULT 2 US-09-810-310-24

Length 15; Indels

Score 77; DB 10; Pred. No. 1.9e-05; ; Mismatches 0;

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APPLICANT: KINSEY, BERNAA
APPLICANT: KINSEY, BERNAA
APPLICANT: KINSEY, BERNAA
APPLICANT: KINSEY, BERNAA
APPLICANT: BHOGAL, BALBIR
TITLE OF INVENTION: MAGENTS
TOTAL OF THE MAGENTS
TOTAL
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Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity 100.0
Matches 15; Conservative
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                                                                             GENERAL INFORMATION:
APPLICANT: Khleif, Samir N.
APPLICANT: Khleif, Samir N.
APPLICANT: BEZZÓFGY, JAY A.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR CO-STIMULATION OF
TITLE OF INVENTION: IMMUNOLOGICAL RESPONSES TO PEPTIDE ANTIGENS
TITLE OF INVENTION: IMMUNOLOGICAL RESPONSES TO PEPTIDE ANTIGENS
TITLE OF INVENTION: IMMUNDER: 15280-415100018
CURRENT FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/189,396
PRIOR APPLICATION NUMBER: 60/189,396
PRIOR APPLICATION NUMBER: 60/189,396
PRIOR SEQ ID NOS: 61
SOFTWARE: PATENTIN VOY: 2.1
SEQ ID NO 24
LENGTH: 15
TYPE: PRT
TYPE: PRT
CORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTHER INFORMATION: Description of Artificial Sequence: HIV PEPTIDE COTHER INFORMATION: ANTIGEN US-09-810-310-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Cheong, Hong-Seok
TITLE OF INVENTION: Liposomes Comprising Peptide Antigens
TITLE OF INVENTION: Derived from X Protein of Hepatitis B virus
FILE REFERENCE: 0136/08154
CURRENT APPLICATION NUMBER: 02/09/989,621
CURRENT FILING DATE: 2001-11-20
PRIOR APPLICATION NUMBER: 09/051,006
PRIOR APPLICATION NUMBER: 09/051,006
SPIOR APPLICATION NUMBER: 09/051,006
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100.0%; Score 77; DB 9; Length 15;
Best Local Similarity 100.0%; Pred. No. 1.9e-05;
Matches 15; Conservative 0; Mismatches 0; Indels
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APPLICANT: Kim, Tae-Youn
APPLICANT: Lee, Ki-Young
APPLICANT: Chang, Jin-Soo
APPLICANT: Chang, Jin-Soo
APPLICANT: Hwang, Yu-Kyeong
APPLICANT: Choi, Myeong
Sequence 24, Application US/09810310
Patent No. US20020044948A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/09989621 Patent No. US20020151683A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIORGPGRAFVTIGK 15
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Best Local Similarity
Matches 15; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 4
US-09-827-688-9
; Sequence 9 %---
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                                                                                                                                                                                                                                                                               Sequence 246, Application US/10133210; Publication No. US20030103964A1; GENERAL INFORMATION: APPLICANT: DeList, Charles
                                                                                                                                                                                                                                             1 RIORGPGRAFVTIGK 15
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; Sequence 9, Application US/09827688 ; Publication No. US20030165476A1

Gaps

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Indels

Length 15;

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Sequence 2, Application US/10787880
; Publication No. US20050025777A1
; GENERAL INFORMATION:
; APPLICANT: Sheehy, Michael J.
; APPLICANT: Sheehy, Michael J.
; APPLICANT: Barton, Kenneth A.
; TITLE OF INVENTION: PARTICLE-MEDIATED DELIVERY OF ANTIGENS
; TITLE OF INVENTION: PARTICLE-MEDIATED DELIVERY OF ANTIGENS
; CURRENT APPLICATION NUMBER: US/10/787,880
; CURRENT FILING DATE: 2004-02-27
; PRIOR APPLICATION NUMBER: US/09/191,772
; PRIOR APPLICATION NUMBER: US/09/191,772
; RIGH FILING DATE: 198-11-13
; NUMBER OF SEQ ID NOS: 2
; SOFTWARE: FASELSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 77; DB 17; Length 15; 100.0%; Pred. No. 1.9e-05;
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i Sequence 44, Application US/10062710

i Publication No. US2000049253A1

i GENERAL INFORMATION:

APPLICANT: Li, Frank Q.

APPLICANT: Chu, Yong-Liang

APPLICANT: Chu, Yong-Liang

TITLE OF INVENTION: Polymeric Conjugates for Delivery of

TITLE OF INVENTION: Wia Peptide Vaccines

TITLE OF INVENTION: Wia Peptide Vaccines

TITLE OF INVENTION: Wia Peptide Vaccines

TITLE OF INVENTION: UNABER: US/10/062,710

CURRENT APPLICATION NUMBER: US 60/310,498

PRIOR FILING DATE: 2001-08-08

NUMBER OF SEQ ID NOS: 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 77; DB 14; 1
Best Local Similarity 100.0%; Pred. No. 1.9e-05;
Matches 15; Conservative 0; Mismatches 0;
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                                                                                                                             TITLE OF INVENTION: Vaccine Composition
FILE REFERENCE: 033267-006
CURRENT FILING DATE: 2002-05-20
PRIOR APPLICATION NUMBER: US 60/291,654
PRIOR PILING DATE: 2001-05-18
PRIOR FILING DATE: 2001-05-18
PRIOR FILING DATE: 2001-05-18
PRIOR FILING DATE: 2001-05-18
NUMBER: OF SEQ ID NOS: 53
SEQ ID NO 6
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Best Local Similarity 100.
Matches 15; Conservative
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; ORGANISM: HIVGD120
US-10-787-880-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: HIV
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US-10-787-880-2
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LENGTH: 15
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'APPLICANT: Berzofsky, Jay

'APPLICANT: Gulukota, Kamalakar

'APPLICANT: Vaccaro, Dennis

APPLICANT: Waccaro, Dennis

APPLICANT: Wang, Zhiping

APPLICANT: Wang, Zhiping

TITLE OF INVENTION: COMPOSITIONS THERBOF

TITLE OF INVENTION: COMPOSITIONS THERBOF

FILE REFERENCE: BU-035AX

CURRENT APPLICATION NUMBER: US/10/133,210

CURRENT APPLICATION NUMBER: US/10/133,210

NUMBER OF SEQ ID NOS: 281

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 246

LENGTH: 15
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; Sequence 262, Application US/10133210
; Publication No. US20030103964A1
; GENERAL INFORMATION:
; APPLICANT: DeLisi, Charles
; APPLICANT: Delisi, Charles
; APPLICANT: Gulukota, Kamalakar
; APPLICANT: Weng, Zhiping
; CURRENT APPLICATION NUMBER: US/10/133,210
; CURRENT FILING DATE: 2002-04-26
; SEQ ID NOS: 281
; SEQ ID NOS: 281
; LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-10-133-210-246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-10-133-210-262
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100.0%; Score 77; DB 14; Length 15;
Best Local Similarity 100.0%; Pred. No. 1.9e-05;
Matches 15; Conservative 0; Mismatches 0; Indels
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; Sequence 6, Application US/10147910
; Publication No. US20030124718A1
; GENERAL INFORMATION:
; APPLICANT: Fuller, Deborah
; APPLICANT: Fuller, James
; APPLICANT: ARynes, Joel
; APPLICANT: Shipley, Timothy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
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Matches 15; Conservative
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Gaps

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FUSION PROTEINS IN A MAMMALIAN CELL
     TITLE OF INVENTION: FUGLOW. ENGLAND FUGLE REPERENCES: ONIO7E/30436.18USD3 FULKER REPERENCES: ONIO7E/30436.18USD3 CURRENT APPLICATION NUMBER: US/10/283,610A CURRENT FILING DATE: 2002-10-29 PRIOR APPLICATION NUMBER: 08/539,436 PRIOR FILING DATE: 1995-10-05 PRIOR PILING DATE: 1995-10-05 PRIOR PILING DATE: 1995-10-05 PRIOR FILING DATE: 1993-00-13 PRIOR FILING DATE: 1993-00-13 PRIOR FILING DATE: 1993-02-01 PRIOR FILING DATE: 1993-02-01 STOWBER OF SEQ ID NOS: 35 SOFTWARE: PATENTIN VEY: 2.1 SEQ ID NO 3 LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Unknown Organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
CRGANISM: Homo sapiens
US-10-283-610A-3
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Best Local Similarity
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US-10-178-488-25
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APPLICANT: Licabetter, Jeffrey A.
APPLICANT: Hayden, Martha S.
APPLICANT: Hayden, Martha S.
APPLICANT: Hayden, Martha S.
APPLICANT: Hayden, Jurgen
APPLICANT: Bajorath, Jurgen
APPLICANT: Gilliland, Lisa K.
TILE OF INVENTION: ENPERSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC
TITLE OF INVENTION: WORDING: US/09/813,659
CURRENT APPLICATION NUMBER: US/09/813,659
CURRENT APPLICATION NUMBER: US/09/49,067
PRIOR FILING DATE: 2000-04-13
PRIOR FILING DATE: 1993-09-13
PRIOR FILING DATE: 1993-02-01
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 3
LENGTH: 20
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Publication No. US20030219876A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ledbetter, Jeffrey A.
APPLICANT: Linsley, Peter S.
APPLICANT: Bajorath, Jurgen
APPLICANT: Fell, H. Perry
APPLICANT: Fell, H. Perry
APPLICANT: Expression vectors encoding bispecific Fusion Proteins
TITLE OF INVENTION: AND METHODS OF PRODUCING BISPECIFIC FUSION PROTEINS
TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC
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                                                                                                                                                        ) OTHER INFORMATION: HIV Helper-T Cell Epitopes US-10-062-710-44
       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 44 LENGTH: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 3, Application US/09813659; Patent No. US20020012989A1; GENERAL INFORMATION:
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                                                                                    TYPE: PRT ORGANISM: Artificial Sequence
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US-09-813-659-3
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Best Local Similarity
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US-09-813-659-3
                                                                                                                                   FEATURE:
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; Sequence 160, Application US/10621675
; Publication No. US2050049398A1
; GENERAL INFORMATION:
; APPLICANT: De Levy. Robert
; TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
; TITLE OF INVENTION: PO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR USE IN
                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Cool Shi-Xian
APPLICANT: Cool, Shi-Xian
APPLICANT: Cool, Shi-Xian
APPLICANT: Person, Roy
APPLICANT: Person, Roy
APPLICANT: Ferson, Roy
APPLICANT: Ferson, Roy
APPLICANT: FIRST PROVINGS-LIKE PARTICLES MADE NON-INFECTIOUS BY A
TITLE OF INVENTION: PLURALITY OF MUTATIONS
TITLE OF INVENTION: PLURALITY OF MUTATIONS
FILE REFERENCE: 1038-1238 MIS
CURRENT FILING DATE: 2002-10-25
PRIOR PRILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 25
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; FEATURE:
; OTHER INFORMATION: Description of Unknown Organism: Artificial
US-10-178-488-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
100.0%; Score 77; DB 15; Length 20; 100.0%; Pred. No. 2.5e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
100.0%; Score 77; DB 14;
Best Local Similarity 100.0%; Pred. No. 2.6e-05;
Matches 15; Conservative 0; Mismatches 0;
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OTHER INFORMATION: Description of Artificial Sequence: Synthetically Designed OTHER INFORMATION: Peptide
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APPLICANT: CORVAIA, Nathalie
APPLICANT: BECK, Alain
APPLICANT: GESTSCH, Lilaine
TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Best Local Similarity 100.0%; Pred. No. 0.00014;
Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 73; DB 16;
Pred. No. 0.00011;
0; Mismatches 1;
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APPLICANT: Li, Frank Q.
APPLICANT: Chi, Yong-Liang
APPLICANT: Chi, Yong-Liang
APPLICANT: Chi, Jian-Tai
TITLE OF INVENTION: Polymeric Conjugates for Del
TITLE OF INVENTION: Wie-Recognized Epitopes
TITLE OF INVENTION: Wis Peptide Vaccines
FILE REFERENCE: 3781-001-27
CURRENT APPLICATION NUMBER: US/10/062,710
CURRENT APPLICATION NUMBER: US/10/062,710
PRIOR APPLICATION NUMBER: US 60/310,498
PRIOR FILING DATE: 2001-08-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , OTHER INFORMATION: HIV Helper-T Cell Epitopes
US-10-062-710-45
                     CURRENT APPLICATION NUMBER: US/10/398,932
CURRENT FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: PCT/JP01/08893
PRIOR PILING DATE: 2001-10-10
PRIOR APPLICATION NUMBER: JP 2000/314288
PRIOR PILING DATE: 2000-10-13
NUMBER OF SEQ ID NOS: 55
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 232
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 45
LENGTH: 18
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Publication No. US20030175285A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
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     FILE REFERENCE: 024918-0103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 93.3
Matches 14; Conservative
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Sequence 1, Application US/10311111
Sequence 1, Application US/10311111
Sequence 1, Application OS US20030121065A1
GENERAL INFORMATION:
APPLICANT: SHIBA, KIYOTAKA
TITLE OF INVENTION:
TITLE OF INVENTION:
CURRENT APPLICATION NUMBER: US/10/311,111
CURRENT APPLICATION NUMBER: US/200-180997
PRIOR FILING DATE: 2000-6-16
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PATENTIN VERSION 3.1
A PROCESS FOR DETERMINATION OF ANTIBODIES OF BIOTIVIZATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT ELITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ô
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Publication No. US20040171803A1
GENERAL INFORMATION:
APPLICANT: SHIBA, KIYOTAKA
APPLICANT: OHNO, TSUNEYA
TITLE OF INVENTION: OP EPITOPE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 24;
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Pred. No. 0.00011;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                              TILLE REPERENCE: 2752-11
CURRENT APPLICATION NUMBER: US/10/621,675
CURRENT PILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: US/09/576,824A
PRIOR APPLICATION NUMBER: US/09/576,824A
PRIOR APPLICATION NUMBER: 08/723,425
PRIOR FILING DATE: 1996-09-30
PRIOR FILING DATE: 1993-11-22
PRIOR FILING DATE: 1993-11-22
PRIOR FILING DATE: 1993-09-08
PRIOR FILING DATE: 1993-03-08
PRIOR FILING DATE: 1992-03-06
PRIOR PRIUNG DATE: 1992-03-06
PRIOR SEQIENTING NUMBER: EP 92400598.6
PRIOR SEQIENTING VALVESTORM NUMBER: EP 92400598.6
PRIOR PRILING VALVESTORM NUMBER: EP 92400598.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Designed peptide US-10-311-111-1
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Best Local Similarity 93.3%;
Matches 14; Conservative
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ORGANISM: artificial
  TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-621-675-160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 160
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RESULT 22
US-09-827-345-24
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US-10-239-313A-186

Sequence 186, Application US/10239313A

Sequence 186, Application US/10239313A

Publication No. US2003017528SA1

GENERAL INFORMATION:
HAMOUR - HAMOUR, Christine
APPLICANT: KINGUER - HAMOUR, Christine
APPLICANT: GORVALA, Nathalie
APPLICANT: GORVALA, Nathalie
APPLICANT: GORSCH, Liliane
APPLICANT: GORSCH, Liliane
APPLICANT: GORSCH, Liliane
APPLICANT: GORSCH, Liliane
TITLE OF INVENTION: NOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION NUMBER: US/10/239,313A

CURRENT APPLICATION NUMBER: PCT 01/70772

PRIOR APPLICATION NUMBER: PCT 01/70772

NUMBER OF SEQ ID NOS: 697

SOPTWARE: ParentIN Ver. 2.1

TENGRALL OF 186
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TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID FILE REFERENCE: 343 727 - US CURRENT APPLICATION NUMBER: US/10/239,313A CURRENT FILING DATE: 2002-09-19 PRIOR APPLICATION NUMBER: FR 00/03711 PRIOR APPLICATION NUMBER: PT 01/70772 PRIOR APPLICATION NUMBER: PCT 01/70772 PRIOR APPLICATION NUMBER: PCT 01/70772 PRIOR PILING DATE: 2000-03-22 NUMBER OF SEQ ID NOS: 697 SOFTWARE: PATCHIL VET: 2.1 SEQ ID NO 536 LENGTH: 13
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                                                                                                                                                                                                                                                                                                                                                                                                                    Length 13;
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100.0%; Pred. No. 0.00044;
ative 0; Mismatches 0;
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US-09-993-307-21
Sequence 21, Application US/09993307
Sequence 21, Application US/09993307
Publication No. US20030162733A1
GENERAL INFORMATION:
APPLICANT: HANTRS, JOG I R.,
APPLICANT: ARRINGTON, JOSHUA
ITILE OF INVENTION: NUCLEIC ACID ADJUVANTS
FILE REFRENCE: APPLICATION NUMBER: US/09/993,307
CURRENT APPLICATION NUMBER: US/09/993,307
CURRENT APPLICATION NUMBER: G0/253,381
                                                                                                                                                                                                                                                                                                              ; TYPE: PRT
; ORGANISM: Human immunodeficiency virus
US-10-239-313A-536
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13; Conservative
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Best Local Similarity
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Matches
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Sequence 535, Application US/10239313A

Sequence 535, Application US/10239313A

Publication No. US20030175285A1

GENERAL INFORMATION:

APPLICANT: CORVAIA, Nathalie

APPLICANT: CORVAIA, Nathalie

APPLICANT: CORVAIA, Nathalie

APPLICANT: GENERAL: Alain

APPLICANT: GENERAL: Alain

TITLE OF INVENTION: McLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS

TITLE OF INVENTION: Wolbecture of A DUTAMIC ACID OR A GLUTAMINE IN THE FORM

TITLE OF INVENTION: Wolber: US/10/239,313A

CURRENT FILING DATE: 2002-09-19

PRIOR FILING DATE: 2000-03-23

PRIOR APPLICATION NUMBER: PT 01/70772

PRIOR APPLICATION NUMBER: PT 01/70772

PRIOR PILING DATE: 2001-03-22

NUMBER OF SEQ ID NOS: 697

SOFTWARE: PATENTIN VET: 2.1

SECTIMARE: PATENTIN VET: 2.1
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APPLICANT: CHERMANN, JEAN-CLAUDE
APPLICANT: GALEA, PASCALE
APPLICANT: GALEA, PASCALE
APPLICANT: GALEA, PASCALE
TITLE OF INVENTION: VACCINE AGAINST INFECTIOUS AGENTS HAVING AN
TITLE OF INVENTION: INTRACELLULAR PHASE, COMPOSITION FOR THE TREATMENT AND
TITLE OF INVENTION: PREVENTION OF HIV INFECTIONS, ANTIBODIES AND METHOD OF
TITLE OF INVENTION: DIAGNOSIS
FILE REFERENCE: 05691-0216
CURRENT APPLICATION NUMBER: 09/599,549
PRIOR APPLICATION NUMBER: 09/599,549
PRIOR APPLICATION NUMBER: PCT/FR96/01006
PRIOR PLING DATE: 1996-06-28
                                                                                                                                                                                                                                                                                                   Gaps '
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100.0%; Pred. No. 0.0025;
ive 0; Mismatches 0; Indels
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PRIOR FILING DATE: 2000-11-27
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin version 3.1
SEQ ID NO 21
LENGTH: 15
                                                                                                                                                                                                                                              85.7%;
86.7%;
                                                                                                                                     ; TYPE: PRT
; ORGANISM: synthetic construct
US-09-993-307-21
                                                                                                                                                                                                                                                                                                                                                                                      1 RIQRGPGRAFVITGK 15
                                                                                                                                                                                                                                                                                                                                                       1 RIQRGPGRAFVTIGK 15
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Best Local Similarity 86.7
Matches 13; Conservative
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Matches 12, Conservative
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; ORGANISM: Homo sapiens
US-10-239-313A-535
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetically Designed
OTHER INFORMATION: Peptide
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/901,106
FILING DATE: 10-Jul-2001
GLASSIFICATION DATA:
APPLICATION DATA:
RILING DATE: 13-AUG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                             Length 13
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                                                                                                                                                                                                                                       1; Indels
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Liljestrom, Peter
TITLE OF INVENTION: DNA Expression Systems Based
Alphaviruses
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ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                           Score 58; DB 16;
Pred. No. 0.017;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Murphy Jr., Gerald M.
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 828-103P
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 10:
US-09-901-106-10
                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 10, Application US/09901106; Patent No. US20020151067A1
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SEQUENCE CHARACTERISTICS:
    SEQ ID NO 3
LENGTH: 13
TYBE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                               75.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 27
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Best Local Similarity 91.7
Matches 11; Conservative
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US-09-901-106-10
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Matches
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Publication No. US20030121065A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: MULTIFUNCTIONAL BASE SEQUENCE AND ARTIFICIAL GENE CONTAINING THE PILLE REPERENCE: 4439-4004
CURRENT PELLOR OF THE 2002-12-13
PRIOR APPLICATION NUMBER: US/10/311,111
CURRENT PILLING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: UP 2000-180997
PRIOR APPLICATION NUMBER: JP 2000-06-16
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin version 3.1
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; Sequence 3, Application US/2039832
; Publication No. US20040171803A1
; GENERAL INFORMATION:
APPLICANT: SHIBA, KIYOTAKA
; APPLICANT: SHIBA, KIYOTAKA
; TITLE OF INVENTION: APPLICALLED PROTEINS WITH ENRICHED IMMUNOCEN
; TITLE OF INVENTION: OF EPITOPE
; FILE REPREBACE: 024918-0103
; CURRENT APPLICATION NUMBER: US/10/398,932
; CURRENT FILING DATE: 2003-04-11
; PRIOR FILING DATE: 2001-10-10
; PRIOR FILING DATE: 2001-10-10
; PRIOR FILING DATE: 2000-10-13
; COMPMANDER OF SEQ ID NOS: 55
; COMPMANDER OF SEQ ID NOS: 55
                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; CTHER INFORMATION: peptide US-09-827-345-24
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Pred. No. 0.017;
0; Mismatches 1; Indels
PRIOR APPLICATION NUMBER: 08/973,551
PRIOR FILING DATE: 1998-02-19
PRIOR PILING DATE: 1998-06-10
PRIOR FILING DATE: 1995-06-30
NUMBER OF SEQ ID NOS: 28
SEQ ID NO 24
LENGTH: 20
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Best Local Similarity 100.
Matches 12; Conservative
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Best Local Similarity 91.7
Matches 11, Conservative
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LENGTH: 13
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TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
Alphaviruses
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                                                                                                                                                     COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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Best Local Similarity 100.
Matches 10; Conservative
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Best Local Similarity
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LENGTH: 10
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                                                                                                       APPLICANT: KLINGUER - HAMOUR, Christine
APPLICANT: CORVAIA, Nathalie
APPLICANT: CORVAIA, Nathalie
APPLICANT: CORVAIA, Nathalie
APPLICANT: CORVAIA, Liliane
TITLE DE INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: MOTERIAL OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: 0.0 F A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: 0.0 F A PHYSTOLOGICALLY ACCEPTABLE STRONG ACID
FRICKRENT FILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: PCT 01/70772
PRIOR APPLICATION NUMBER: PCT 01/70772
PRIOR PELLING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 697
SEQ ID NO 533
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; Sequence 6, Application US/10622003
; Publication No. US20050014230A1
; GENERAL INFORMATION:
; APPLICANT: Chin, Li-Te
; TITLE OF INVENTION: PREPARATION OF FULLY HUMAN ANTIBODIES
; FILE REFERENCE: 16863-002001
; CURRENT APPLICATION NUMBER: US/10/622,003
; CURRENT FILING DATE: 2003-07-16
; NUMBER OF SEQ ID NOS: 20
; SOFTMARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 6
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100.0%; Pred. No. 0.029;
tive 0; Mismatches 0; Indels
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TITLE OF INVENTION: DNA Expression Systems Based on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Synthetically generated peptide
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                                         Sequence 533, Application US/10239313A Publication No. US20030175285A1 GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 11; Conservative
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ORGANISM: Mus musculus
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Matches 11; Conserv
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                        US-10-239-313A-533
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US-09-901-106-12
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Sequence 3, Application US/09858349
Patent No. US20020012909A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: SMALL FUNCTIONAL UNITS OF ANTIBODY HEAVY CHAIN VARIABLE REGIONS
FILE REFERENCE: 87534-2800
CURRENT APPLICATION NUMBER: US/09/858,349
CURRENT FILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.1
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100.0%; Pred. No. 0.12;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              74.0%; Score 57; DB 9;
100.0%; Pred. No. 0.032;
iive 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/920,281C
FILING DATE: 13-AUG-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Murphy Jr., Gerald M.
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 828-103P
                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/901,106
FILING DATE: 10-Jul-2001
CLASSIFICATION: <unknown>
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US-09-087-513-13
JS-09-087-513-7
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             Sequence 16, Application US/09810310
| Sequence 16, Application US/09810310
| Patent No. US20020044948A1
| GENERAL INFORMATION:
| APPLICANT: Khleif, Samir N. APPLICANT: Khleif, Samir N. APPLICANT: Berzofsky, Jay A. TITLE OF INVENTION: IMPUNOLOGICAL RESPONSES TO PEPTIDE ANTIGENS
| TITLE OF INVENTION: IMPUNOLOGICAL RESPONSES TO PEPTIDE ANTIGENS
| TITLE OF INVENTION: IMPUNOLOGICAL RESPONSES TO PEPTIDE ANTIGENS
| TITLE OF INVENTION: IMPUNOLOGICAL RESPONSES TO PEPTIDE ANTIGENS
| FILE REFERENCE: 15280-415100US
| CURRENT APPLICATION NUMBER: 06/189,396
| PRIOR APPLICATION NUMBER: 60/189,396
| PRIOR APPLICATION NUMBER: 60/189,396
| NUMBER OF SEQ ID NOS: 61
| SOFTWARE: Patentin Ver. 2.1
| SOFTWARE: Patentin Ver. 2.1
| SOFTWARE: PRIOR APPLICATION NUMBER: PRIOR APPLICATION NOT NOT TYPE: PRIOR APPLICATION NOT TYPE: 
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OTHER INFORMATION: Description of Artificial Sequence: HIV PEPTIDE
OTHER INFORMATION: ANTIGEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  67.5%; Score 52; DB 9;
100.0%; Pred. No. 0.12;
tive 0; Mismatches
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Best Local Similarity 100.0%; Pred. No. 0.12;
Matches 10; Conservative 0; Mismatches
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Patent No. US20020142977A1
GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 10; Conservative
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US-09-810-310-16
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Sequence 7, Application US/09087513;
Publication No. US20020182180A1;
GENERAL INFORMATION:
APPLICANT: KOZBOR, Danuta
TITLE OF INVENTION: METHOD OF INDUCING IMMUNITY TO VIRUSES
FILE REFERENCE: 0010-0929-0X;
CURRENT APPLICATION WUMBER: US/09/087,513
CURRENT FILING DATE: 1998-05-29
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.1
LENGTH: 10
                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence:peptide US-09-087-513-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: KANEKO, YULAZO
APPLICANT: KOZBOR, Danuta
TITLE OF INVENTION: METHOD OF INDUCING IMMUNITY TO VIRUSES
FILE REFERENCE: 0010-0529-0X
CURRENT APPLICATION NUMBER: US/09/087,513
CURRENT FILING DATE: 1998-05-29
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 13
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APPLICANT: Bowers, William J.
APPLICANT: Bowers, William J.
APPLICANT: Frelinger, John G.
APPLICANT: Willis, Richard A.
APPLICANT: Evans, Thomas J.
APPLICANT: Toba, Khaled A.
APPLICANT: Toba, Khaled A.
APPLICANT: Rosenblatt, Joseph D.
TITLE OF INVENTION: HELPER VIRUS-FREE HERPESVIRUS AMPLICON
                                                                                                                                                                                                                                                                                                                                                                                       Length 10;
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Pred. No. 0.12;
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Pred. No. 0.12;
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100.0%; Pred. No. v.
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ORGANISM: Human immunodeficiency virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-087-513-13; Sequence 13, Application US/09087513; Sequence 13, Publication No. US20020182180A1; GENERAL INFORMATION:
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100.0%;
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Best Local Similarity 100.
Matches 10; Conservative
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Best Local Similarity 100.
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Query Match 67.5%; Score 52; DB 14; Length 10; Best Local Similarity 100.0%; Pred. No. 0.12; Matches 10; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.
Matches 10, Conservative
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       ; ORGANISM: Homo sapiens
US-10-113-085-7
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Best Local Similarity
Matches 10; Conserva
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Publication No. US20030003086A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Rock, Kenneth L.
APPLICANT: Goldberg, Alfred L.
TITLE OF INVENTION: MODULATION OF WHC CLASS 1 ANTIGEN PRESENTATION
FILER REFERENCE: 07917-140001.
CURRENT APPLICATION NUMBER: US 10/113,085
CURRENT FILING DATE: 2002-09-04
PRIOR APPLICATION NUMBER: US 60/280,669
NUMBER OF SEQ ID NOS: 12.
SOFTWARE: FASELSEQ for Windows Version 4.0
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TITLE OF INVENTION: PARTICLES AND USES THEREOF

CURRENT APPLICATION NUMBER: US/09/997, 848A

CURRENT FILING DATE: 2002-09-10

PRIOR PELICATION NUMBER: US 60/253, 858

PRIOR PILING DATE: 2000-11-29

PRIOR PILING DATE: 2000-11-30

PRIOR PILING DATE: 2000-11-30

NUMBER OF SEQ ID NOS: 18

SOFTWARE: FREESEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-993-307-22

Sequence 22, Application US/09993307

Sequence 22, Application US/09993307

GENERAL INFORMATION:

APPLICANT: HAYNES, Joel R.

APPLICANT: ARRINGTON, JOSHUA

TITLE OF INVENTION: NUCLEIC ACID ADJUVANTS

FILE REFERENCE: APP41.20

CURRENT FILING DATE: 2001-11-26

PRIOR FILING DATE: 2000-11-27

PRIOR PILING DATE: 2000-11-27

NUMBER OF SEQ ID NOS: 26

SOFTWARE PATENTIN VUMBER: 60/253,381

FILOR FILING DATE: 2000-11-27

NUMBER OF SEQ ID NOS: 26

SOFTWARE PATENTIN VUMBER: 60/253,381

FILOR FILING DATE: 2000-11-27

NUMBER OF SEQ ID NOS: 26

SEQ ID NO 22

LENGTH: 10
                                                                                                                                                                                                                                                                                                                   ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; .TYPE: PRT; ORGANISM: synthetic construct
US-09-993-307-22
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Best Local Similarity 100.
Matches 10; Conservative
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                                                                                                                                                                                                                                                SEQ ID NO 16
                                                                                                                                                                                                                                                                                               TYPE: PRT
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Sequence 2, Application US/10168843A

Sequence 2, Application US/10168843A

Publication No. US20030108562A1

GENERAL INFORMATION:

APPLICANT: Medical Research Council

APPLICANT: University of Nairobi

TITLE OF INVENTION: Improvements in or Relating to Immune Responses to HIV

FILE REFERENCE: MJL/C124811/M

CURRENT FILING DATE: 2002-09-24

NUMBER OF SEQ ID NOS: 33

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 2

SEQ ID NO 2

SEQ ID NO 3.
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100.0%; Pred. No. 0.12;
tive 0; Mismatches 0; Indels
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APPLICANT: Fuller, James
APPLICANT: Haynes, Joel
APPLICANT: Haynes, Joel
APPLICANT: Shipley, Timothy
TITLE OF INVENTION: Vaccine Composition
FILE REFERENCE: 032567-006
CURRENT APPLICATION NUMBER: US 60/291,654
PRIOR APPLICATION NUMBER: US 60/291,655
PRIOR APPLICATION NUMBER: US 60/291,655
PRIOR PILING DATE: 2001-05-18
PRIOR PILING DATE: 2001-05-18
NUMBER OF SEQ ID NOS: 53
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12
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Gaps

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Query Match 67.5%; Score 52; DB 14; Length 10; Best Local Similarity 100.0%; Pred. No. 0.12; Matches 10; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 0.12;
tive 0; Mismatches 0; Indels
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APPLICANT: Raz, Eyal

APPLICANT: Rai, Hearn Jay

APPLICANT: Richman, Douglae

APPLICANT: Richman, Douglae

APPLICANT: Horner, Anthony A.

TITLE OF INVENTION: Lymphocyte Response in vivo.

FILE REPERRACE: UCAL-188D1V.

PRIOR FILING DATE: 2000-03-28

PRIOR PILING DATE: 2000-03-28

PRIOR PILING DATE: 2000-05-11

PRIOR PILING DATE: 2000-05-11

PRIOR PILING DATE: 2000-07-05

NUMBER OF SEQ ID NOS: 8

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO
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| Publication No. US20030147870A1
| GENERAL INFORMATION:
| APPLICANT: Raz, Eyal
| APPLICANT: Rac, Hearn Jay
| APPLICANT: Richman, Douglass
| APPLICANT: Richman, Douglass
| APPLICANT: Richman, Lymphoryte Response in vivo.
| TITLE OF INVENTION: Method for Increasing a Cytotoxic T
| TITLE OF INVENTION: Method for Increasing a Cytotoxic T
| TITLE OF INVENTION: Method for Increasing a Cytotoxic T
| TITLE OF INVENTION: Method for Increasing a Cytotoxic T
| TITLE OF INVENTION: Method for Increasing a Cytotoxic T
| TITLE OF INVENTION: Method for Increasing a Cytotoxic T
| TITLE OF INVENTION: Method for Increasing a Cytotoxic T
| FILE REFERENCE: UCAL-188CON
| CURRENT APPLICATION NUMBER: US/10/339,885
| PRIOR APPLICATION NUMBER: US 60/192,537
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US-10-340-275-8
; TYPE: PRT
; ORGANISM: Unknown
; FRATURE:
. OTHER INFORMATION: CTL Peptide Epitope of HIV gag
US-10-079-167-68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 8, Application US/10340275; Publication No. US20030143213A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 10; Conservative
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1 RGPGRAFVTI 10
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US-10-079-167-51
                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: McShane, Helen
APPLICANT: McShane, Helen
APPLICANT: Galbert, Sarah C.
APPLICANT: Reece, William
APPLICANT: Schneider, Joerg
TITLE OF INVENTION: Vaccination Method
TITLE REFERENCE: 2907.1000-001
CURRENT APPLICATION NUMBER: US/10/079,167
CURRENT FILING DATE: 1999-12-09
PRIOR PILING DATE: 1999-12-09
PRIOR APPLICATION NUMBER: GB 97 11957.2
PRIOR APPLICATION NUMBER: GB 97 11957.2
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 2001-09-13
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APPLICANT: Millin, Adrian V.S.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Schneider, Joerg
TITLE OF INVENTION: Vaccination Method
FILE REPERBINE: 2007.1000-001
CURRENT APPLICATION NUMBER: US 09/454,204
PRIOR APPLICATION NUMBER: US 09/454,204
PRIOR PILING DATE: 1999-12-09
PRIOR PLING DATE: 1999-12-09
PRIOR FILING DATE: 1999-06-09
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 2001-09-13
PRIOR FILING DATE: 2001-09-13
PRIOR FILING DATE: 2001-09-13
PRIOR FILING DATE: 2001-09-13
PRIOR FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 99
SCOTTWARR: FASHESEQ for Windows Version 4.0
SEQ ID NO 68
LENGTH: 10
                                                                                           US-10-079-167-51

Sequence 51, Application US/10079167

Publication No. US20030138454A1

GENERAL INFORMATION:
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1 RGPGRAFVTI 10
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RESULT 45
US-10-360-836-48
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TITLE OF INVENTION: Compositions And Methods For Modulation Of Immune Response
FILE REFERENCE: TROMOROZ.
CURRENT APPLICATION NUMBER: US/10/210,148
FILOM PAPLICATION NUMBER: POT/US02/24311
PRIOR PAPLICATION NUMBER: POT/US02/24311
PRIOR FILING DATE: 2002-07-31
NUMBER OF SEQ ID NOS: 117
SOFTWARE: Patentin version 3.2
LENGTH: 10
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US-10-206-155-5
i Sequence 5, Application US/10206155
j Publication No. US2003015/135A1
j Publication No. US2003015/135A1
j APPLICANT: TSUJi, Moriya
j APPLICANT: TSUJi, Moriya
j APPLICANT: Nussenzweig, Ruth S.
j APPLICANT: Nussenzweig, Ruth S.
j APPLICANT: Nussenzweig, Ruth S.
j APPLICANT: Norsulka, Yasuliko
j TITLE OF INVENTION: USE OF GLYCOSYLCERAMIDES AS ADJUVANTS
j TITLE OF INVENTION: FOR VACCINES AGAINST INFECTIONS AND CANCER
j FILE REPERENCE: 5986/14958US1
CURRENT APPLICATION NUMBER: US/10/206,155
j RUOR APPLICATION NUMBER: 2002-07-25
j RUOR PILING DATE: 2001-07-25
j NUMBER OF SEQ ID NOS: 19
j SOPTWARE: FRSESEQ for Windows Version 3.0
j SEQ ID NO 5
j LENGTH: 10
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                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: HIV-1 class I-restricted gp120 peptide
US-10-339-885-8
PRIOR FILING DATE: 2000-03-28
PRIOR APPLICATION NUMBER: US 60/203,567
PRIOR FILING DATE: 2000-05-11
PRIOR PILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 10; Conservative
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; ORGANISM: HIV-1
US-10-206-155-5
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TYPE: PRT
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GENERAL INCORDITION: USE OF Fidel
APPLICANT: Birkett, Ashley
TITLE OF INVENTION: USE OF RECOMBINANT HEPATITIS B CORE
TITLE OF INVENTION: PARTICLES TO DEVELOP VACCINES AGAINST INFECTIOUS PATHOGENS
TITLE OF INVENTION: PARTICLES
FILE REFERENCE: 5986/1J876
CURRENT PAPLICATION NUMBER: 0210/360,836
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/354,963
RIOR FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 86
SOFTWARE: FASTERQ for Windows Version 3.0
SEQ ID NO 48
LENGTH: 10
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                                                                                        Score 52; DB 14; Length 10;
Pred. No. 0.12;
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                                                                                                                                       0; Indels
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; ORGANISM: human immunodeficiency virus (HIV-1)
US-10-360-836-48
                                                                Query Match 67.5%; Score 52; DB
Best Local Similarity 100.0%; Pred. No. 0.1
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nes 10; Conservative
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                   ; ORGANISM: Homo sapiens
US-10-210-148-113
; TYPE: PRT
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Aaw72821 HIV-1 gp1 Aaw8620 Epitope o Aay04680 HIV-1 gp1 Aay83916 HIV-1 env Aay66439 HIA-A2-bi Aay66455 HIA-A3-bi Aay85591 HIV relat Aab15875 Human che Aab15875 Human che Aab92348 Virus rel	HIV 9F Human HIV ef HIV-1 HIV-1 HIV CT HIV DE	MHC by Altere Altere Altere Altere Lipope HIV-D HIV-1 HIV-1 HIV-1 HIV-1 HIV-1 HIV-1	Aav16052 HIV-1 180 Aaw85568 Human imm Aab15012 Peptide P Aau08699 Retroviru Aax42153 990120 V3 Aax57470 HIV BRU V Aaw07392 HIV-1 Btr Aay07488 HIV-1 Btr Aay85137 HIV-1 Btr Aay85137 HIV-1 Btr Aay86137 HIV-1 N-1	CCpd. e HIV.1 HIV.1 Secarce Secarce Secarce HIV GF Secarce HIV GF Secarce HIV.1
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ion 5.1.6 05 Compugen Ltd. lel Search time 165 Seconds (without alignments) 35.160 Million cell updates/s		768190	s predicted by chance to have a score of the result being printed, total score distribution.  ARIES  Description	Aap82095 Env-KI pe Aap91228 Peptide c Aar21143 HIV-1 gpl Aar31207 Sequence Aar51619 V3 loop p Aar51619 V3 loop r Aar68189 Cytotoxic Aaw68189 Cytotoxic Aaw68535 HIV-1 gpl Aar92007 HIV-1 gpl Aaw92031 HYdrophil Aaw07311 gpl20 pep Aaw32010 HIV-1 wy Aaw07311 gpl20 pep Aaw3201 HIV-1 wy Aaw42010 CA+ T-1y Aaw10348 HIV epito Aaw32031 Antigenic Aaw32031 Antigenic Aaw32031 HIV-1 syn Aaw4036 HIV-1 syn Aaw4036 Fushon im Aaw5689 Sequence Aay26896 Sequence Aay26896 Sequence Aay26896 Sequence Aay26896 HIV gpl20 Aay26896 Sequence Aay26896 HIV gpl20 Aay26896 Sequence

Aar04060 Epitope c Aae20149 Human imm Aar63820 HIV-1 gpl Aaw22329 HIV-1 cli Aaw62892 Peptide s Aar62152 HIV-1 gpl20 Aam99432 HIV gpl20 Aam99432 Vaccine r Aaw76864 Fusion imm Aar04476 Human imm Aar04476 Human imm	Aar1219 V3 peptid Aar31278 HIV princ Aar26714 HIV princ Aar26601 Alkaline Aar58601 Alkaline Aar58602 Alkaline Aar31254 HV princ Aar31254 HV princ	Aar26689 HIV-PND-p Aar56689 HIV-PND-p Aar3336 Sequence Aar48604 Sequence Aar6167 HIV-1 Str Aar62167 HIV-1 Gpl Aar6852 Fusion im Abb05777 HIV gpl20 Aao15659 Strong im Aax24218 CD4+ T-1y Aar32408 Sequence Aar68799 Cytotoxic	Amm99430 Vaccine r Abp17102 HIV B27 s Aar10592 Protease Aar16302 HIV-1 gp1 Aav76846 Pusion im Adp76013 Peptide e Aar2513 SFV-HIV e Aar2513 SFV-HIV e Aar67313 HIV-1 pep Aaw67331 HIV-1 pep Aaw9939 HIV-1 pep Aaw9939 HIV-1 Wac Aay9690 HIV-1 Wac Aay9690 HIV-1 Mar91029 Aar6618 HIV-1 III Aar90229 Cyclic HI Aar62165 HIV-1 Gp1	Aav76861 Fusion im Aav76861 Hru princ Aar04441 Human imm Aar6865 T cell ep Aav25835 Hru B-cel Aav89959 Hrv 1 vac Aav89959 Hrv 1 vac Aav89757 Hru Chim Aav22593 Hru cpito Aar26890 Hrv epito Aar26892 Hrv epito Aar26892 Hrv epito Aar26892 Hrv epito Aar36899 Hrv epito Aar36899 Hrv epito Aar36899 Hrv epito Aar36899 Hrv epito Aav76899 Fusion im Aav10172 T cell ep Aav1057 Hrv Cell ep Aav10547 Hrv Gass Aav10164 T cell ep Aav10164 T cell ep Aav10165993 Hrv gag C Aav93655 Hrv aag C Aav93659 Hrv age C Aav96559 Hrv gag C Aav96559 Hrv gag C Aav96559 Hrv gag C Aav96589 Hrv gag C Aav96589 Hrv gag C Aav96589 Hrv pepti
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25 2 AAR30031 Aar30031 25 2 AAR36712 Aar3322 25 2 AAR41336 Aar41336 25 2 AAR41330 Aar41336 25 2 AAR41330 Aar41330 25 2 AAR41330 Aar41330 25 2 AAR972819 Aar72819 25 2 AAR97413 Aar72819 25 4 AAR09522 Aar86419	15 2 AAR66430 Aar66430 20 5 ABB05775 Abb05775 20 5 AAO15657 Aao15657 14 2 AAR66416 Aar66416 15 1 AAP95357 Aar66416 15 2 AAR33460 Aar533460 15 2 AAR62166 Aar633460 15 2 AAR66427 Aar66428 15 2 AAR66427 Aar66428	15   2 AAR66426   Aar66426     16   2 AAR33236   Aar33236     17   1 AAP95349   Aap955349     17   1 AAP95349   Aar29241     17   2 AAR29241   Aar22241     17   2 AAR69664   Aar68664     17   2 AAR75834   Aar68664     17   2 AAR75834   Aar76848     17   2 AAR75839   Aar76848     17   2 AAR75850   Aar76848     17   2 AAR75850   Aar76848     17   2 AAR799558   Aar76849558     18   2 AAR799558   Aar76845     19   2 AAR799558   Aar76845     10   2 AAR799558   Aar76845     11   2 AAR799558   Aar76845     12   AAR799558   Aar76845     13   AAR799558   Aar76845     14   Aar76845     15   AAR76845     16   Aar76845     17   AAR76845     18   Aar76845     18   Aar76845     19   Aar76845     10   Aar76845     11   AAR76845     12   AAR76845     13   AAR76845     14   Aar76845     15   Aar76845     16   Aar76845     17   Aar76845     18   Aar76845     18   Aar76845     18   Aar76845     18   Aar76845     19   Aar76845     10   Aar76845     11   Aar76845     12   AAR76845     13   Aar76845     14   Aar76845     15   Aar76845     16   Aar76845     17   Aar76845     18	17 7 ADM14075 Adm1407	88.3 13 2 AAM62890 Peptide E88.3 13 2 AAM62891 Peptide E88.3 13 4 AAM99433 Aam99433 Vaccine r Aam69433 Vaccine r Aam66417 Aaw76897 Pusion in E88.3 18 2 AAM63062 Aam99083 Vaccine r Aam99091 Vaccine r Aam99091 Vaccine r Aam99091 Vaccine r Aam99997 Vaccine r Aam99999 Vall vaccine r Aam9999 Vall vaccine r Aam99999 Vall vaccine r Aam9999 Vall vaccine r Aam99999 Vall vaccine r Aam9999 Vall vaccine r Aam99999 Vall vaccine r Aam99999 Vall vaccine r Aam9999 Vall vaccine

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Chimeric immunoglobulin; viral-neutralising; HIV-1;
BAT123 mouse immunoglobulin; viral antigen-binding region; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                        Peptide will induce an immune response in subject, and will thus act as on infective vaccine, prophylactic or have therapuetic value for AIDS patients. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 27-AUG-2003 to correct OS field.) (Updated on 24-OCT-2003 to standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV-1 gp120 epitope found in mouse immunoglobulin BAT123 and mouse/human chimeric antibody CAG1-51-4.
                                                                                                                                                                                                                                                                                                                                     Synthetic human immuno-deficiency virus env-coded peptide(s) - induce antibodies that block human immuno-deficiency virus proliferation and fusion between infected and non-infected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 77; DB 1; Length 15; 100.0%; Pred. No. 9e-05; tive 0; Mismatches 0; Indels
                                                                     Peptide comprising AAs 308-322 of HIV-I IIIB env protein
                                                                                                                                                                                                                                                     PONT DE NEMOURS & CO E I.
PONT MERCK PHARMACEUTICAL CO.
                                                                                                                                                                                                                                                                                         Durda PJ;
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                                                                                                                      Human immunodeficiency virus 1
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89US-00324027.
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                                                                                              AIDS; HIV-I; vaccine
                                                                                                                                                                                                                                                                                                               WPI; 1989-317386/44.
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 AA;
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                                                                                                                                                                                            21-APR-1989;
                       27-AUG-2003
25-MAR-2003
                                                                                                                                                                                                                  6-APR-1988;
                                                                                                                                                                                                                             20-MAR-1989;
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16-MAY-1992
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                                                                                                                                            EP339504-A.
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  g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This peptide elicits cytotoxicity by T-cells against HIV antigens and stimulates prodn. of HIV-specific cytotoxic T-lymphocytes (CTLs). It is specific for the HIV envelope protein gp160. (Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent clashes with ongoing US granted patent numbers. For further information please visit the Derwent web site at www.derwent.com/dwpj/updates/ntis us.html.) (Updated on 25-MAR-2003 to correct PP field.) (Updated on 25-MAR-2003 to correct PP field.)
Aab15874 Human che
Aab92350 Virus rel
Aab43397 HIV pepti
Aae04801 Human imm
Aae20153 Human imm
Abg31255 GP120 cla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic peptide corresp. to HIV GP 160 ENV sequence - which elicits cytotoxicity by T cells against HIV and proliferation of HIV-specific cytotoxic T cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Moss B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 1; Length 15; 100.0%; Pred. No. 9e-05; artive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                            Env-K1; gp160 Env protein; T-cell cytotoxicity; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Germain R,
                                                                                                 ALIGNMENTS
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 AAB15874
AAB92350
AAB49397
AAE04801
AAE20153
ABG31255
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAP91228 standard; peptide; 15 AA
                                                                                                                                                          AAP82095 standard; peptide; 15 AA
                                                                                                                                                                                                                                                                                                                                                                            88US-00148692
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(first entry)
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  22222
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Best Local Similarity
Matches 15; Conserv
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17-DEC-2001
29-OCT-1990
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                                                                                                                                                                                                                                                                                                       Synthetic.
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2449
250
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Gaps

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Chang NT;

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The inventors claim a chimeric, viral-neutralising immunoglobulin which binds to the gp120 region of HIV-1 with a potency and immunologic specificity equal to BA7123 mouse Ig. It comprises a viral-specific antigen-binding region of non-human origian and a constant region of non-human origian and a constant region of the non-human origian and a constant region of a 47439 percent of a 47430 p
                                                                                                                                                                                                                New chimeric HIV-1-neutralising immunoglobulin(s) - comprising non-human antigen binding regions and constant human region, for immuno-therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gp120; HIV-1; cytotoxic T-lymphocyte; CTL; T-helper; AIDS; infection.
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                                                                                                                                Chang TW,
                                                                                                                                Fung MS,
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                                                                                                                                                                                                                                                                                                         Example; Page 26; 39pp; English
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90WO-US004048
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                                                                                                                                Rosen EM, Sun BN,
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                                                                                      (TANO-) TANOX BIOSYST INC
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nes 15; Conser
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18-JUL-1990;
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12-OCT-1993
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HIV gpl20 V3 loop-derived peptides (AAR38170-87) are successful in generating CTL responses, esp. peptide R15K (AAR38187); the T-helper cell-inducing peptide includes the sequence C19A (AAR38164); HIV infection-inhibiting peptides are given in AAR38188-206, and are esp. peptides R15K, N24G, E13V, R8K, T13Q and H13N (AAR38165-69). The peptides may also be derived from an influenca virus protein or a sendai virus protein (AAR41014-15). It was observed that peptide R15K (amino acids 315-329), with sequences derived from the V3 loop of HIV-1 IIIB, inhibits HIV-1 infection of primary human T cells by 92* at 1 microg/ml (ca. 0.4-0.6 microw). (Updated on 25-MAR-2003 to correct DN field.) (Updated on 27-AUG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The peptide corresponds to amino acid residues numbered 315-329 in the V3 loop of the envelope glycoprotein gpl60 of human immunodeficiency virus (HIV-1), as numbered by Ratner in the strain MN. It is useful for the prophylaxis or immunotherapy of HIV-1 infection. It elicits an immunised
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    73 loop; envelope glycoprotein; gp160; HIV-1; prophylaxis; immunotherapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 puence of peptide which corresp.to AA residues 315-329 of the V3 loop the gpl60 envelope glycoprotein in {\rm HIV-1} strain {\rm MN.}
                                                                   Peptide composition for treating and preventing viral infections -
comprise CTL-inducing epitope and HIV infection-inhibiting sequence or
helper cell-inducing sequence.
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 Nebete PN;
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Sastry JK, Arlinghaus RB, Platsoucas CD,
                                                                                                                                       Claim 13 + 19; Page 94-95; 130pp; English.
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17-DEC-2001
07-JUN-1993
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AAR74603 standard; peptide; 15 AA.
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                                                AAR74603;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New fusion polypeptide of antigen binding domain and HIV epitope - useful as vaccine for treatment or prevention of HIV infection, ensures efficient focusing of epitopes on surface of antigen presenting cells.
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subject ctytoxic T lymphocyte (CTL) activity against the corresp. cilnical isolate of HIV-1. (Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent clashes with ongoing US granted patent numbers. For further information plases wist the Derwent web site at www.derwent.com/dwpi/updates/ntis_us.html.) (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                     gp 120; HIV epitope; Human Immunodeficiency Virus fusion polypeptide.
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                                                                                                                                                                                                                                                                                                                                               V3 loop region of gp120 of HIV.
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hes 15; Conservative
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21-OCT-1994
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MAD 5023 was developed against AA residue 308-322 of the variable loop of HIV-I (ARR74663). MAD 5023 binds to the epitope GRAF. G preceding RAF is believed to critical for internalization. MAD 5023 recognised p160, p120, p42 and p24 in cancer cells. ARR74603 competitively blocked binding of the MAD to the cancer antigens, indicating that at least the epitope GRAF, which is recognised by the MAD, must also be present in cancer antigens. (Updated on 25-MAR.2003 to correct PI field.) (Updated on 15-MAR.2003 to correct PI field.) standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detection of HIV-1 cross-reactive breast carcinoma-associated antigens for diagnosis and anti:sense therapy of breast and gynaecological
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                                                                                                                                                      MAb 5023; variable V3 loop; HIV-I; human immunodeficiency virus; cancer antigen; monoclonal antibody.
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                                                                                                           HIV-I variable loop residues 308-322
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                                                                                                                                                                                                   Synthetic peptides spanning multideterminant regions from the HIV envelope protein gp160 have been designed and are designated cluster peptides (PCLUS). These peptides each consist of a cluster of overlapping determinants and are known to induce in vitro T cell proliferation and cytokine production in mice and humans of multiple MHC types. The cluster peptides were co-linearly synthesised at the N-terminus of an immunodominant CTL determinant, Peptide 18 (AAR66414), corresp. to part of the gp160 V3 loop and principal neutralising determinant region of HIV - IIIB isolate. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytotoxic I lymphocyte; epitope; antigen; pathogenic; nef; gag; pol; env; gpl20; gp41; HIV; cell-mediated immunity; human immunodeficiency virus; class I restricted.
                                                                                                                                              Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
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Best Local Similarity 100.
Matches 15, Conservative
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23-AUG-1995
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                                       24-NOV-1994
  Synthetic.
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AAR68789
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                                                                                                                                                               AAR68744-805 are cytotoxic T lymphocyte (CTL) class I and II restricted epitopes derived from human immunodeficiency virus proteins. AAR68789 corresponds to amino acid residues 308-322 of the env gpl20 protein. These antigens are examples of peptides that can be used with an immunogenic implant. The implant is associated with an antigen associated with a pathogen and used to form a discrete, localised chronic inflammation site which acts as a local 'factory' for prodn. of CTL's pathogen-specific CTL's can eradicate or prevent development of infection, and can also be used to treat or arrest the development of cancers associated with infection. (Updated on 25-MAR-2003 to correct PN
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      forming
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Implant for sustained release of pathogen-associated antigen - form chronic inflammatory site producing cytotoxic T-lymphocytes lysing infected cells, esp. for treating AIDS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 77; DB 2; Length 15;
Pred. No. 9e-05;
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                                                                                                                       Disclosure; Page 12; 35pp; English
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 AA;
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Determining the likelihood of maternal transmission of HIV-1 to foetus by measuring maternal reactivity with specific gp120 and gp41 derived peptide(s), also used for diagnosing HIV in infants.
                                                                                                                                                                                                                          HIV; gpl20; HIV-IIIB strain; HIV-1 transmission; foetal transmission; neutralising antibody; passive immunisation; anti-idiotypic antibody; gp41; vaccine; active immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYPE-) UNIV PENNSYLVANIA.
(WIST-) WISTAR INST ANATOMY & BIOLOGY.
                       AAW07931 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ugen KE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Col 18; 63pp; English,
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                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          94US-00218025.
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                                                                                                                 (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Williams WV, Weiner DB,
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es 15; Conserva
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                                                                                                                                                                                gp120 peptide p18p.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-MAR-1994;
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                                                                                                              16-OCT-2003
31-JAN-1997
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                                                                    AAW07931;
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AAR92007
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polypeptides contain a first portion of 5-99 amino acids joined to a second portion contg. at least a functional domain of epimorphin. The first portion may be selected from the peptides given in AAR92039 to AAR92036. The second portion may be full-length epimorphin (see AAR92037 to AAR92042 for human and mouse epimorphins). Fragments of epimorphins given in AAT16091 to AAT16090 are used in the prodm. of modified epimorphins are useful for the development of diagnosis and treatment of morphogenetic abnormalities of epithelial tissue or novel remedies for wounds, eg burns, after surgery and for artificial organs. They may also be used as components of cosmetics, hair growth stimulators, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bpimorphin, human, mouse, wound; burn; epithelial tissue, diagnosis,
treatment; morphogenetic abnormality; cosmetic; hair growth stimulator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel polypeptide containing an epimorphin functional domain - has possible benefits in epithelial tissue treatments, e.g. burns and for
                                                                                                                 Gaps
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                                                                    100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 9e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hydrophilic peptide for epimorphin modification (5).
                                                                                                                                                                                                                                                                                                                      AAR92033 standard; peptide; 15 AA.
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                                                                                                                                                                                                     1 RIORGPGRAFVTICK 15
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                                                                    Query Match
Best Local Similarity 100.0
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-1996 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1996-118213/13.
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   artificial organs.
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                            Sequence 15 AA;
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31-MAR-1995;
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                                                                                                                                                                                                                                                                          RESULT 11
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AAR9

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AAR9

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This sequence represents a HIV gpl20 peptide that can be used in the method of the invention is for determining whether or not a mother will transmit HIV-1 to a foetus. The method comprises incubating a sample from the HIV-1 infected mother, with a comprises incubating a sample from the HIV-1 infected mother, with a collection of HIV peptides includes at least one of the gpl20 sequences (such as AAMO7909-W07917), and at least one HIV gp41 derived peptide (see AAMO7918-W0793). The number of peptides that react with the sample is determined, and this number is compared with a candard that shows pattern reactivity for a patient of fransmission status. A non-transmissive and this sundleds if the test sample can be presence of neutralising antibodies that protect against mother to infant transmission of HIV. These sequences can also be used in vaccines to protect against transmission. Antibodies against these sequences can be used for passive immunisation, and to generate anti-idiotypic antibodies for use in vaccines or active immunotherapy. (Updated on 16-correct correct against mother to active immunotherapy. (Updated on 16-correct correct sequences or active immunotherapy. (Updated on 16-correct correct c
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RESULT 12

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Antigenic peptides can be inserted into the alpha-antigen sequence of a Mycobacterium and secreted from an appropriately transformed M.bovis BCG cell. The resulting chimeric antigen has greatly enhanced antigenicity and immunogenicity and is recognised in vivo by B-cells which recognise the alpha-antigen. The present sequence is that of a HIV-1 gpl20 V3 loop epitope which was incorporated into the alpha antigen. M. bovis BCG cells secreting a chimeric protein comprising the epitope sequence are useful as anti-AIDS vaccines. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                             BCG containing vaccine secretes chimeric protein containing foreign antigen - has enhanced immunogenicity and antigenicity esp. when used as an anti-AIDS vaccine.
                                                        Mycobacterium bovis BCG; AIDS vaccine; surface protein; alpha antigen;
Human immunodeficiency virus type 1; fusion protein; gp120 epitope;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CD4+ T-lymphocyte epitope to HIV-1 V3 loop derived peptide V3-LAI-P18.
                           HIV-1 V3 loop epitope, for insertion in Mycobacterium alpha antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         T-lymphocyte epitope; diagnosis; antigen; infectious disease; delayed-type hypersensitivity assay; vaccine development.
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                                                                                                                                                                                                                                                                                                                      Yamazaki S,
                                                                                                                                                                                                                                                                                                                      Yamazaki A, Honda M,
                                                                                                                                                                                                                                                                           (AJIN ) AJINOMOTO CO INC.
(NINA-) JAPAN AGENCY NAT INST HEALTH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW24219 standard; peptide; 15 AA
                                                                                                                     Human immunodeficiency virus 1.
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27-SEP-1996 (first entry)
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                                                                                                                                                                                                                                                                                                                        Matsuo K, Chujo Y,
                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-129127/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 AA;
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                                                                                          V3 loop.
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AAW24219
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the T-lymphocyte epitopes of a large antigent to be determined in vivo in humans. The method is useful in medicine e.g. in diagnosis, monitoring and treatment design for infectious disease exposure, active autoimmune disease. Allergic diseases and mallipancy. It is especially useful for tracing infectious diseases and mallipancy. It is especially useful for tracing infectious diseases e.g HIV, particularly when a sequence is present only in certain strains of an organism, and developing suitable vaccines. Vaccinated individuals can also be tested to verify protection against a particular strain. The method allows in vivo mapping of T-lymphocyte epitopes, not previously possible. The method is simpler, more rapid and more sensitive. It can also be applied in a variety of environments e.g. undeveloped regions since specialist equipment is not
                                                                                                                                                                                                                                                                                                             Peptide fragments AAW24217-20 were used to demonstrate a new method of tracing sources of infectious diseases. The method comprises preparing a short (9-50 amino acid) peptide containing at least one non-conserved epitope of an organism, injecting a composition containing the peptide intradermally into a test subject in a delayed-type hypersensitivity (DTH) assay and observing the infection site at intervals for induration. In this example CD4+ T-lymphocyte epitopes to the HIV-1 V3 loop were mapped by existing in vitro technique for two existing HIV infected individuals and used to design peptides AAWA2217-20. The method allows
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                                                                                                                                                                      Peptide fragments containing antigen epitope(s) used to trace diseases used in a delayed-type hypersensitivity assay, for in vivo mapping of human T-lymphocyte epitope(s) e.g. for diagnosis, vaccine development
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 77; DB 2; Length 15; 100.0%; Pred, No. 9e-05; ive 0; Mismatches 0; Indels
                                                      (USSA ) US DEPT ARMY GOVERNMENT US ARMY MEDICAL.
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                                                                                                                                                                                                                                                                                Disclosure, Page 6, 14pp, English.
                96US-0010679P.
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Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIQRGPGRAFVTIGK
                                                                                                                                    WPI; 1997-393814/36
                                                                                                 Sitz KV, Brix DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15 AA;
                  26-JAN-1996;
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Tasaka H;

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(USSH ) US DEPT HEALTH & HUMAN SERVICES

96WO-US010108 95US-00488435

07-JUN-1996; 07-JUN-1995;

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20-FEB-1998 (first entry
                                                                                                  Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                              (HARD ) HARVARD COLLEGE
                                                                                                                                                                                                                                                WPI; 1997-350782/32.
           WPI; 1997-108658/10.
                                                                                       Sequence 15 AA;
                                                                                seroconversion
                                                                                                                                                                                                                                         Starnbach MN;
                                                                                                                                                                                                            13-DEC-1996;
                                                                                                                                                                                                                   13-DEC-1995;
                                                                                                                                                                                             WO9723236-A1
                                                                                                                                                                                                                      07-JUN-1996;
                                                                                                                                                                                                     03-JUL-1997
   Shearer GM,
                                                                                                                                              AAW22031;
                                                                                              Query Match
                                                                                                                                RESULT 16
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This is the antigenic human immunodeficiency virus peptide P18. This antigenic compound can be introduced into the cytoplasm of a cell by a contacted with a fusion molecule comprising a delivery molecule. The delivery molecule can either be a polycationic affinity handle, LFN (the protective antigen binding domain of anthrax compound is linked to either of the delivery molecules by a covalent compound is linked to either of the delivery of the antigenic compound is linked to either of the delivery of the antigenic compound of an operated fusion proteins with a toxin B molety, which alleviates the need to generate fusion proteins with a toxin B molety, which alleviates problems associated with incorrect folding of lengthy fusion proteins. Small cationic fusion peptides substituted for LFN may reduce the spossibility of sterric interference with the biological activity of the translocated protein. The method is used for the introduction of antigens, e.g. MHC class I antigens or any other therapeutic protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptides AAW39275-W39300 are used in a novel method for inducing cytotoxic T-cell (CTC) activity specific for a broad array of HIV-1 isolates using hybrid synthetic peptides. The method involves first administering a recombinant viral vector expressing the HIV-1 gp160 envelope glycoprotein and then administering at least 1 chimeric synthetic polypeptide. When several synthetic polypeptides having sequences corresponding to amino acids 315-329 of the gp160 envelope glycoprotein of HIV-1 strain IIIB, in which amino acid 325 is substituted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Inducing cytotoxic T-cell response to HIV - by administering {\tt gpl60} vector and chimeric {\tt gpl60} peptide(8).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus type I; HIV-1; cytotoxic T-cell; CTC; vaccine; prophylactic; immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 9e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Germain RN, Berzofsky JA, Takahashi H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW39275 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Col 3; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93US-00095332.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88US-00148692.
91US-00760530.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1 synthetic peptide IIIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                         cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JAN-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-SEP-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                         into the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW39275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                       The present sequence represents a synthetic HIV-1 gp160 peptide env P18-IIB for use in a method for diagnosing exposure of a patient to human immunodeficiency virus (HIV). The method involves: (a) obtaining peripheral blood mononuclear cells (BPMC) from a patient; (b) incubating the PBMC with at least 1 synthetic peptide representing an epitope(s) of the infectious agent (e.g. the present sequence); and (c) determining the activation of the PBMC as a result of the incubation in step (b). The method can provide for the early detection of exposure to infectious organisms, specifically HIV in this case. The method can be used to assess exposure to HIV without concomitant infection. It also provides an earlier identification of HIV exposure, than is provided by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antigenic peptide; human papillomavirus; MAGB gene; BAGB-1 peptide; P18; human immunodeficiency virus; cancer antigen; tyrosinase; signal procein; anthrax lethal factor; LF; toxin; cationic fusion peptide; translocation; gene therapy; polycationic affinity handle; therapeutic protein; LFN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Introducing therapeutic proteins, especially antigens, into cells - using toxin molecules and/or polycationic handles for delivery.
                                                                                                                          Diagnosis of exposure to infectious agents, partic. HIV - by detecting activation of peripheral blood mononuclear cells from patient by epitope of infectious agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 77; DB 2; Length 15;
Pred. No. 9e-05;
; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antigenic human immunodeficiency virus peptide P18.
                                   Clerici M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW22031 standard; peptide; 15 AA.
                                                                                                                                                                                                                                Claim 15; Page 62; 82pp; English.
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                                      Berzofsky JA,
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Gaps

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0; Indels

(first entry)

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B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                       Fusion immunoglobulin heavy chain HIV gp120 T cell epitope #17.
             AAW76898 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                              (AMNA-) AMERICAN NAT RED CROSS.
                                                                                                                                                                                                                                                                                                                                                                          Scott D, Zambidis E;
                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-506315/43.
                                                                                                                                                                                                                                           WO9836087-A1
                                                                                                                                                                                                                                                                                              13-FEB-1998;
                                                                                                                                                                                                                                                                                                                       13-FEB-1997;
                                                                                                                                                                                                                  Нощо варіепв
                                                                  25-JAN-1999
                                                                                                                                                                                                                                                                    20-AUG-1998.
                                         AAW76898;
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AAW76898
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a fragment of the human immunodeficiency virus (HIV) Type 1 strain IIIB gp120 protein. This sequence is used in a method resulting in the production of a vaccine against HIV which induces an immune response to the V3 epitope of HIV. This method which comprises the transplantation of an epitope of HIV at plural sites in the hypervariable region of immunoglobulin, the preparation of the epitope molecule histocompatibility antibody, and optionally chemically cross linking the epitope. An epitope histocompatibility antibody is also described in the specification which specifically responds to HIV, prepared by transplantation of an epitope comprising a peptide obtained from at least one V3 sequence of HIV. (Updated on 17-0CT-2003 to standardise OS field)
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by the corresponding amino acid from other strains or isolates, are used, a CTC response to a broad range of HIV-1 isolates can be elicited. These synthetic peptides are useful as vaccines for the prophylaxis or immunotherapy of HIV-1 virus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vaccine against human immunodeficiency virus - induces immune response reaction to V3 epitope of virus.
                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                              Epitope, vaccine, V3, gpl20, immune response, hypervariable region; immunoglobulin, histocompatability antibody.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              th 100.0%; Score 77; DB 2; Length 15; Similarity 100.0%; Pred. No. 9e-05; 15; Conservative 0; Mismatches 0; Indels
                                                                                             Score 77; DB 2; Length 15;
Pred. No. 9e-05;
; Mismatches 0; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                AAW40316 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                        HIV-1 IIIB gp120 peptide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 5; 8pp; Japanese.
                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1.
                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96JP-00232378
                                                                                                                                                    1 RIORGPGRAFVTIGK 15
                                                                                                                                                                              RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                                           Ouery Match
Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                 17-OCT-2003
23-JUN-1998
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97US-0040581P.

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This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human common than fused in frame at its N-terminus to one or more human immunoglobulin have the first positopes. Pusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune disease. Induction of tolerance suppresses production of antibodies against gpl20, so prevents or inhibits 'bystander' apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral continuantly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
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                                                                                                                                                                        Claim 11; Page 120; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW54929 standard; peptide; 15 AA.
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tes 15, Conservative
                                                                                                            virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 AA;
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ID AAW5
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1 RIORGPGRAFVTIGK 15 RIORGPGRAFVTIGK 15

Local Similarity

Matches

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Gaps

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The invention relates to vaccine compositions comprising a vaccine and a timed-release formulation of a cytokine or cytokine/lmmunoglobulin fusion protein or plasmid. The formulation or device releases the cytokine protein or plasmid at one or more temporal points subsequent to vaccine administration. The vaccines can be used for treating an autoimmune disease, an inflammatory disease, a neoplastic disease, an inflammatory disease, a neoplastic disease, or an immunologic disease in an individual. The vaccines can be used to elicit immune responses against disease such as ALDS, malaria, tuberculosis, hepatitis C, hepatitis B, cancer or influenza. The methods can provide for enhancement of one or more immunologic parameters such as an antibody response, a cellular proliferative response as well as cytotoxic T-lymphocyte levels. In addition the Ig can increase the circulating half life of the cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes peptide antigens AAY24459 to AAY24463 derived from the X protein of hepatitis B virus (HBV) which are recognized by cytocoxic T lymphocytes (CTL). The peptide antigens derived from HBV X protein are useful for inducing CTLs against the virus or inducing immunological tolerance to the virus. pH-sensitive liposomes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis B virus, HBV; X protein; cytotoxic T lymphocyte; liposome; CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis B virus protein X-derived peptide antigens used to stimulate cytotoxic T lymphocytes, useful for treatment of HBV-associated diseases,
                          New vaccine compositions for treating AIDS, maleria, tuberculosis, cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cheong H;
                                                                                                                                                                                                                                                                                                                                        100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 9e-05; cive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hwang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MOGA-) MOGAM BIOTECHNOLOGY RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 14; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY24466 standard; peptide; 15 AA.
                                                                        Example 3; Page 22; 66pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cho S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antigen; immunity; liver cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98WO-KR000010.
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                                                                                                                                                                                                                                                                                                                                                                                                    1 RIORGPGRAFVTIGK 15
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        especially liver cancer.
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WPI; 1999-254931/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV peptide R15K-1
                                                                                                                                                                                                                                                                                                             Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JAN-1998;
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                                              or influenza
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY24466;
                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kim T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY24466
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                                                                                                                                                                                                                                                                                                                                                                                                                               셤
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                                                                                                                                                                                                                                                                                                                                                                                                                 Peptides AAW54903-W54941 can be used as an immunoadsorbent in an immunoassay for detecting antibodies to HIV gpl20, or as an immunogen for eliciting antibodies to HIV in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fusion protein; vaccine; cytokine; immunoglobulin; autoimmune disease; infectious disease; inflammatory disease; neoplastic disease; cancer; immunologic disease; immune response; malaria; tuberculosis; hepatitis; AIDS; influenza; interleukin; IL-2; Ig.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                              Immunoadsorbent; immunoassay; HIV gp120; immunogen; antibody; Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                           gp120 peptides - useful as immunoassay reagents or vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 9e-05; ive 0; Mismatches 0; Indels
HIV gp120 envelope protein, peptide 127, analogue 127g'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BETH-) BETH ISRAEL DEACONESS MEDICAL CENT.
                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Column 21/22; 34pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence of gp120IIIB P18 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY06896 standard; peptide; 15 AA
                                                                                                                                                                                           91US-00663262.
91US-00726605.
94US-00326676.
                                                                                                                                                                                                                                                      (UNBI-) UNITED BIOMEDICAL INC.
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97US-00990180.
                                                                                                                                                   95US-00488252.
                                                                                                                                                                              88US-00155321.
                                                            Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 RIORGPGRAFVTIGK 15
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Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                WPI; 1998-347301/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-SEP-1998;
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                                                                                                                                                07-JUN-1995;
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                                                                                         US5763160-A
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                                                                                                                                                                                                                                                                                                                                                             components.
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                                                                                                                                                                                                                                                                                     Wang
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Gaps

Sequence 15 AA;

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containing the peptide antigens are used to induce cellular immunity that CTLs specific to the virus can be produced. This is useful for prevention and treatment of HBV-associated diseases, especially HBV-associated liver cancer. pH-sensitive liposomes permit the selective transportation of anti-cancer drugs. The present sequence represents beptide used in an example from the present invention
                                                                                                                                                                                         Sequence 15 AA;
                                                                                                                                                                                                                                         Query Match
             8888888
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100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 9e-05; cive 0; Mismatches 0; Indels 1 RIORGPGRAFVTIGK 15 1 RIORGPGRAFVTIGK 15 Conservative Local Similarity nes 15; Conserv Matches à

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Gaps

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AAY25189 standard; peptide; 15 AAY25189; RESULT 23

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HIV protein gp160 peptide fragment #1. 03-SEP-1999 (first entry)

Heat shock protein; HSP; complex; denatured protein matrix; antigen; vaccine; allergic disease; treatment; susceptibility; Th2; skin rash; allergic reaction; asthma; gp160.

Human immunodeficiency virus.

WO9929182-A1

17-JUN-1999.

98WO-US025734.

97US-00985548. 05-DEC-1997; 05-DEC-1997; (UYNE-) UNIV NEW MEXICO STATE

Moseley PL; Wallen ES,

WPI; 1999-394912/33

Synthesizing heat shock protein complexes using a denatured protein matrix.

Example 1; Fig 1A; 33pp; English.

This invention describes a novel method for synthesizing heat shock protein (HSP) complexes comprises adding a heat shock protein to a denatured protein nearth for binding, and adding a complexing solution comprising a peptide to elute a heat shock protein-peptide complex. A HSP cantigen complex is useful as a vaccine for treating an allergic disease.

C(in a mammal, preferably a human) to reduce susceptibility of the Th2 response, the complex comprising a HSP-antigenic peptide complex. The complex is administered to prevent a mammal from having an allergic reaction to an allergic disease, or administered to a mammal having an allergic disease, to reduce the allergic reactions. Allergic diseases include asthm and skin reaches which react only the symptoms, corricosteroids which have severe side effects and desensitization thereby which has limited uses. The new method also allows more flexibility of use of peptide-based vaccines, as prior art HSP-based vaccines require isolation from a portion of the tumour itself. This sequence represents a peptide fragment from the HIV gp160 protein which is used in the method of the invention 

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                                                                                                                                                   Heat shock protein; HSP; complex; denatured protein matrix; antigen; vaccine; allergic disease; treatment; susceptibility; Th2; skin rash; allergic reaction; asthma; V3 protein.
                     Gaps
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      DB 2; Length 15;
100.0%; Score 77; DB 2; Lenguarion 100.0%; Pred. No. 9e-05;
                                                                                          AAY25204 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                        (UYNE-) UNIV NEW MEXICO STATE.
                                                                                                                                                                                                                                                        97US-00986234.
                                                                                                                                                                                   Human immunodeficiency virus.
                                                                                                                                                                                                                                 98WO-US025734.
                                                                                                                                                                                                                                                97US-00985548.
                                                                                                                                       HIV V3 peptide fragment #10.
                                     1 RIQRGPGRAFVTIGK 15
                                                   1 RIORGPGRAFVTIGK 15
                                                                                                                        (first entry)
       Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                       Wallen ES, Moseley PL;
                                                                                                                                                                                                                                                                                                     WPI; 1999-394912/33
                                                                                                                                                                                                  WO9929182-A1
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                                                                                                                        03-SEP-1999
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                                                                                                                                                                                                                  17-JUN-1999.
                                                                                                        AAY25204;
                                                                           RESULT 24
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This invention describes a novel method for synthesizing heat shock protein (HSP) complexes comprises adding a heat shock protein to a denaturated protein matrix for binding, and adding a complexing solution comprising a peptide to elute a heat shock protein-peptide complex. A HSP comprising a peptide to elute a heat shock protein-peptide complex. A HSP complex is useful as a vaccine for treating an allergic disease (in a mammal, preferably a human) to reduce susceptibility of the Th2 response, the complex comprising a HSP-antigenic peptide complex. The complex is administered to a mammal having an allergic complex is administered to a mammal having an allergic disease, to reduce the allergic reactions. Allergic diseases include asthma and skin rashes. Prior art methods or preventing/treating include asthma and skin rashes. Prior art methods or preventing/treating allergic diseases include antihistamines which treat only the symptoms, corticosteroids which have severe side effects and desensitization therapy which has limited uses. The new method also allows more learning regolation from a portion of the tumour itself. This sequence represents a peptide fragment from the HIV V3 protein which is used in the method of the invention

Synthesizing heat shock protein complexes using a denatured protein

Example 1; Fig 1B; 33pp; English.

natrix.

Sequence 15 AA;

Gaps ö 100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 9e-05; tive 0; Mismatches 0; Indels 15; Conservative Best\_Local Similarity Matches 15; Conserv Query Match

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1 RIORGPGRAFVTIGK 15

AAY05356;

AAY05356

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The present invention describes an antibody conjugate comprising an antibody (Ab) which binds to an epitope within amino acid residue 308-322 of gp120 and neutralises HIV-1, conjugated with a cytotoxic agent, an anti-viral agent or an agent which facilitates passage through the blood brain barrier. Also described is an antibody conjugate as above but where the Ab binds to an epitope within amino acid residue 298-312 of gp12 which neutralises HIV-1. The present sequence represents an HIV-1 gp120 monoclonal antibody BAT121 residue 308 to 322 from an example of the protein on the envelope of HIV-1. They inhibit the infection of T-cells and also inhibit syncytium formation. The antibodies are group specific and also inhibit syncytium formation. The antibodies are group specific and also inhibit syncytium formation. The antibodies are group specific and also inhibit syncytium formation. The antibodies are group specific and also thibit syncytium formation. The antibodies are group specific and also thibit syncytium formation the reatment and prevention of AIDS and AIDS related complex. They are especially used to kill infected T-cells. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Epitope; gp120 protein; monoclonal antibody; HIV-1; antibody BAT123;
antibody BAT267; antibody BAT085; T cell infection inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  epitope within amino acid residue of gp120 which neutralises HIV-1 conjugated with, e.g. cytotoxic agent.
                                                                                     HIV-1; gpl20; epitope; monoclonal antibody; envelope; neutralise; inhibit; infection; T-cell; inhibit syncytium formation; AlDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibody conjugate comprising monoclonal antibody - which binds
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                                            HIV-1 gp120 monoclonal antibody BAT123 residue 308 to 322.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW87620 standard; peptide; 15 AA.
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                                                                                                                                                   Human immunodeficiency virus 1.
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89US-00343540.
92US-00895197.
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                                                                                                                                                                                                                                                                                                                            87US-00057445.
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    (first entry)
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Les 15; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 AA;
                                                                                                                                                                                                                                                                                   04-MAR-1993;
      13-JAN-1999
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25-APR-1989;
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03-MAR-1999
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                                                                                                                                                                                              US5834599-A
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                                                                                                                                                                                                                                                                                                            HIV-1; CLUVAC; cluster peptide vaccine construct; cytotoxic T lymphocyte; protective mucosal CTL response; hepatitis A virus; papilloma virus; feline immunodeficiency virus; feline leukaemia virus; M. tuberculosis; Listeria monocytogenes; M. leprae; Giardia lamblia; immune response induction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence represents a HIV-1 cluster peptide vaccine conjugate (CLUVAC) sequence. The invention relates to a method for inducing a protective mucosal cytocaic Tlymphocyte (CTL) response in a mammalian subject, which comprises contacting a mucosal tissue of the subject with induce a protective mucosal CTL response in a subject. The method can be used for protection against e.g., hepatitis A virus, papilloma virus, feline immunodeficiency virus, feline leukaemia virus, insteria monocytogenes, M. tuberculosis, M. leprae, or Giardia lamblia. The method induces long-lasting protective mucosal immune responses. (Updated on 17-ocr-2003 to standardise OS filed)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Method for inducing a protective mucosal cytotoxic T lymphocyte immune
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Derby MA, Kelsall BL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (USSH ) US DEPT HEALTH & HUMAN SERVICE.
                                                                                                                                                                                                                                                                                   HIV-1 CLUVAC peptide, SEQ ID NO. 15.
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                                                                                                                                  AAY05356 standard; peptide; 15 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; Page 85; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98WO-US019028
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1 RIORGPGRAFVTIGK 15
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(first entry)
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29-JUN-1999
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response.

AAW72821;

RESULT 26

AAW72821

Query Match Matches

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Gaps

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The present sequence represents an epitope of the gp120 protein of human immune deficiency virus (HIV)-1. The sequence comprises amino acids 308 to 322 of gp120. The specification describes monoclonal antibodies which bind to epitopes of the gp120 protein. Specifically, these antibodies are designated BAT123, 267 and 085. Monoclonal antibodies neutralise HIV-1, inhibiting both infection of T cells and formation of syncytia, so are complex, by passive immune deficiency syndrome (AIDS) and AIDS-related complex, and in extracorporeal systems. They can also be used as antiviral agents, and in extracorporeal systems. They can also be used as development for neutralising epitopes, potentially useful in vaccine development. (Updated on 20-MMR-2003 to correct PR field.) (Updated on 17-occr-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                            Monoclonal antibodies directed against regions of gp120 of human immune deficiency virus-1 - are neutralisating and able to inhibit infection of T cells and formation of syncytia, used for treatment, prevention or diagnosis of acquired immune deficiency syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gpl20; HIV-1; monoclonal antibody; homology; antigen; breast; prostate;
gynecological; cancer; detection; diagnosis; cell membrane; chromatin.
syncytia formation; acquired immune deficiency syndrome; AIDS; AIDS-related complex; passive immunisation; antiviral; cytotoxic; viral load measurement; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 9e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                 Chang TW;
                                                                                                                                                                                                                                                                                               Chang NT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY04680 standard; peptide; 15 AA.
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                                                                     Human immunodeficiency virus 1.
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87US-00137861.
91US-00767533.
                                                                                                                                                                                                                                                                                                  Sun CRY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Col 8; 16pp; English.
                                                                                                                                                                     92US-00950571.
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(first entry)
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hes 15; Conservative
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                                                                                                                                                                                                                                                                                                    Sun BNC,
                                                                                                                                                                                                                                                                    (TANO-) TANOX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 AA;
                                                                                                                                                                      22-SEP-1992;
                                                                                                                                                                                                                    24-DEC-1987;
26-SEP-1991;
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22-JUN-1999
                                                                                                                                                                                                     29-MAY-1987;
                                                                                                       US5854400-A
                                                                                                                                      29-DEC-1998
                                                                                                                                                                                                                                                                                                  Fung MSC,
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                                                                                                                                                                                                                    This peptide corresponds to amino acids 308-322 from the gpl20 protein of the human immunodeficiency virus type 1 (HIV-1). The peptide is used to generate the monoclonal antibody was 5023. The invention relates to the use of homology between HIV-1 antigens and breast, gynecological and prostate cancer antigens to develop agents for use in the detection and treatment of such cancers. The method especially uses an antibody which recognises the p160, p80, p45 and p24 cell membrane proteins and the p24 chromatin protein. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                  Use of HIV-1 polypeptides - for developing products for the detection and treatment of breast, gynecological and prostate cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Assessing an immune response against a selected agent in an individual comprises accelerating a particulate composition, containing an immunogenic compound from a selected agent, into the target skin site of the individual.
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                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 2; Length 15; 100.0%; Pred. No. 9e-05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                               Disclosure, Page 39; 80pp; English
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                                                                  RAKO/) RAKOWICZ-SZULCZYNSKA E
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            98WO-US015580
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Les 15; Conservative
                                                                                               Rakowicz-Szulczynska EM,
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10-JUN-1999;
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25-FEB-1999.

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                                The invention relates to a method of using an immunogenic compound from a selected agent in the manufacture of a particulate composition for assessance and minume response against the selected agent in an individual. The method comprises: (a) accelerating the particulate composition into a target skin site in the individual; and (b) assessing the target site to determine the presence or absence of a localized skin immune reaction, where the presence of the immune reaction is indicative of an immune response against the selected agent. Peptides AAY83316-Y8392s represent examples of peptides that could be used if the method is used to detect human immunoceficiency virus type 1 (HIV-1). The method is useful for assessing immunocompetence, antibody and cell mediated immunity, antigen exposure, or allergic conditions in an individual. (Updated on 12-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV-1; MHC; major histocompatibility complex; Class I; HLA-A2; human leukocyte antigen; CTL; cytotoxic T-cell; epitope; allele; binding; conserved; genome; peptide; targetting; toxic; drug; antibody; antigen; antiviral; molecular conjugate therapeutic; diagnosis; treatment; pathogen; localisation; quantification; detection; infection; detection; drug resistance; immune response.
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Disclosure, Page 23; 41pp; English.
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than administering it systemically, which may result in adverse side effects. The invention relates to improved methods for the design of molecular conjugate therapeutics for the diagnosis and treatment of infections caused by pathogens with a high mutation rate (such as HIV-1). This method involves identifying conserved peptide-encoding regions among the genomes of multiple variants of a pathogen, identifying the Class I MFC molecules in a population of interest (e.g., human sub-populations), and determining which of the peptides bind to the Class I MFC molecules. The MHC-binding peptides and the construct targetting antigens, which are slatched and used to construct targetting antigens, which are in turn used to produce targetting antibodies. The methods may be used in localisation, complexes and also to detect and treat viral infection. The methods of cupy and also to detect and treat viral infection. The methods of the invention mitigate against the development of viral resistance to drugs and to the immune response, as well as providing a solution for targetting toxic compounds to destroy viruses sequestered in sites not accessible to T cells. In addition, the methods eliminate the virus, whereas current therapies only arrest viral replication. (Updated on 12-SEP-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HLA-A3-binding HIV-1 GP120 CTL epitope #257.
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Best Local Similarity
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cc histocompatibility complex) Class I alleles found with high frequency among Caucasoids in the USA. The peptides are used to construct targetting antigens comprising one or more peptides bound to the corresponding MHC Class I molecule, which can be used to raise antibodies. The antibody may then be used as a targetting vehicle to deliver a potentially toxic drug to its target site of action, rather than administering it systemically, which may result in adverse side effects. The invention relates to improved methods for the design of confections caused by pathogens with a high mutation rate (such as HIV-1). This method involves identifying of a pathogen, identifying the Class I will method involves identifying of a pathogen, identifying the Class I confected by the peptide encoding regions among the genomes of multiple variants of a pathogen, identifying the Class I conferent (e.g., human sub-populations), and determining which of the peptides bind to the Class I MHC molecules. The MHC-binding peptides and the construct targetting antigens, which are in turn used to produce targetting antipodies. The methods may be used in localisation, complexes and also to detect and treat viral infection. The methods of the invention mitigate against the development of viral resistance to drugs and to the immune response, as well as providing a solution for targetting toxic compounds to destroy viruses sequestered in aites not accessible to T cells. In addition, the methods eliminate the virus, whereas current therapies only arrest viral replication. (Updated on 12-cossible to grandardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antiarthritic; antiinflammatory; cell-mediated immune response; cancer; rheumatoid arthritis; inflammatory disorder; viral infection.
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(AGEN-) AGENE RES INST CO LTD.
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Best Local Similarity
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New peptide 3, amide and heterocyclic compounds and saccharide conjugates used for inhibiting chemokine induced activity and for treating e.g. stroke, vascular diseases, autoimmune diseases and tumor growth.
                                      This invention relates to artificial immunogenic particles comprising glycolipids having a lipidic and a saccharide portion and peptide-lipid conjugates having a lipidic and a peptide portion embedded into a lipid based matrix. The peptide portion of the particle may be of viral origin. Peptides AAV85579-Y85591 are human immunodeficiency virus (HIV) related peptides which can be used as the peptide portion in an immunogenic and antihifiammatory activity. The immunogenic particles are used for inducing a cell-mediated immune response in a host directed towards the peptide portion of the peptide-lipid conjugate. This means that the particles may be used to treat diseases such as cancer, rhumatoid arthritis, inflammatory disorders or viral infections such as HIV. (Updated on 12-SEP-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                      100.0%; Score 77; DB 3; Length 15; 100.0%; Pred. No. 9e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human chemokine derived peptide #27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Fig 18; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB15875 standard; peptide; 15 AA.
             Claim 10; Page 34; 39pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-00229071.
99US-00271192.
99US-00452406.
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Best Local Similarity 100...
Best Local Similarity المربقة
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                                                                                                                                                                                                                                                                                                           Sequence 15 AA;
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17-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 33
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exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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                                                                                                                                                                                     AAB92348
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                                                                                                                                 셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidy) and maleimido groups) attached to reactive group (II) (e.g. succinimidy) and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently conds with amino/hydroxyl/thoi groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids.

(I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body.

Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half intercellular uptake and interference with physiological processes.

AAB90829 to AAB92441 represent peptides which can be used in the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity.
autoimmune diseases, atherosclerosis, osteoporosis, HIV infection and AIDS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotoxaemia, myocardial infarction, acute ischaemia and theumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in gene therapy for the same diseases, as well as in the production of animal models
                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                    Protection, endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention describes a modified therapeutic peptide (I)
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                                                                                                                                 Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 702; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                           Virus related peptide SEQ ID NO:1521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Milner PG,
                                                                                                                                                                                                                                                                                AAB92345 standard; peptide; 15 AA.
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99US-0159783P.
                                                                                                                                                                                      1 RIORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                   22-JUN-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-112059/12.
                                                                                                                                   Query Match
Best Local Similarity
                                                                                                            Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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The present invention describes a modified therapeutic peptide (I)

comprising a therapeutically active amino acid region (III) and a
reactive group (II) (e.g. succinimidyl and maleimido groups) attached to
a less therapeutically active amino acid region (IV), which covalently

compared with amino/hydroxyl/thiol groups on blood components to form a
conds with amino/hydroxyl/thiol groups on blood components to form a
condition for modifying therapeutic peptides e.g. hormones, growth
can neurotransmitters, to protect them from peptidase activity in
cators and neurotransmitters, to protect them from peptidase activity in
convo for the treatment of various disorders. Endogenous therapeutic
peptides are not suitable as drug candidates as they require frequent
administration due to rapid degradation by peptidases in the body.
Modifying and attaching therapeutic peptides to albumin prevents or
reduces the action of peptidases to increase length of activity (half
contineracellular uptake and interference with physiological processes.
AAB90829 to AAB92441 represent preptides which can be used in the
compared to the present invention
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                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protection, endogenous therapeutic peptide, peptidase, conjugation, blood component, modification, succinimidyl, maleimido group, amino, hydroxyl; thiol; hormone, growth factor; neurotransmitter.
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                                                   100.0%; Score 77; DB 4; Length 15; 100.0%; Pred. No. 9e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Milner PG, Holmes DL, Thibaudeau
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9e-05;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Virus related peptide SEQ ID NO:1524.
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                                                                                                                                                                                                                                                                                                                                                                                                                            AAB92348 standard; peptide; 15 AA.
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100.0%;
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99US-0153406P.
99US-0159783P.
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                                                                                                                                                                                                                                                          1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                            Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bridon DP, Ezrin AM,
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Sequence 15 AA;
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The invention relates to a composition comprising an expression vector bound to an aggregated protein-polycationic polymer conjugate or suspension. The expression vector contains a promoter polymucleotide sequence operatively linked to a polymucleotide sequence encoding an antigen which is a fragment of a gene or genome associated with an infectious disease, cancer and autoimmune disease such as rheumatoid arthritis, vasculitis, and multiple sclerosis, pathogenic genomes consisting of bacterium, fungus, protozoa and virus such as human immunodeficiency virus (HIV), herpes simplex virus (HSV), hepatitis C virus (HCV), influenza and respiratory syncytial virus (RSV), and optionally comprising a nucleotide sequence encoding a cytokine (or a cytokine expression vector), is useful for inducing an immune response (systemic and/or mucosal) in an organism. The cytokine expression vector contains a sequence for granulocyte macrophage-colony stimulating factor (GM-CSF) or interleukin-12 (IL-12). The polymucleotide sequences encoding the antigen and the cytokine are under transcriptional control of same or
                                                                                                                                                                      HIV; human immunodeficiency virus; cytostatic; immunosuppressive; p18; virucide; antibacterial; fungicide; protozoacide; antirheumatic; vaccine; antiinflammatory; antiarthritic; neuroprotective; rheumatoid arthritis; cancer; multiple soleroais; immune response; vasotropic; gene therapy; autoimmune disease; vasculitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                different promoter polynucleotide sequences. The expression vector, as a DNA vaccine is useful for treating a condition in an organism. The present sequence is human immunodeficiency virus (HIV) p18 peptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Composition for oral delivery of vaccines, comprises expression vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        containing antigenic genomic sequence, bound to aggregated protein-
polycationic polymer conjugate or suspension.
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                                                                                                                                    Human immunodeficiency virus (HIV) p18 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3xample 10, Page 30, 145pp, English.
                      AAE15743 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BAYU ) BAYLOR COLLEGE MEDICINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                07-APR-2000; 2000US-0195680P.
                                                                                                                                                                                                                                                                                            Human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                            06-APR-2001; 2001WO-US011372,
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                                                                                              (first entry)
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1es 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-066308/09.
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                                                                                                                                                                                                                                                                                                                                  WO200176643-A1.
                                                                                                26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                        18-OCT-2001.
                                                          AAE15743;
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    AAE15743
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inposomes having an outer surface layer of polyecthies continued in the content of the distal end. A polypeptide or polyeaccharide effector and cach having a free distal end. A polypeptide or polyeaccharide effector molecule is covalently attached to a portion of the distal ends. The effector interferes with specific binding of pathogen or cell in a bloodstream to a target cell or cell matrix, and is rapidly removed by remain clearance from the bloodstream when administered in free form. The liposome composition may be used in treating a condition mediated by binding a pathogen or cell in the bloodstream, to a target cell or cell matrix. It can be used in treating septic shock, toxic shock, colonic matrix. It can be used in treating septic shock, toxic shock, colonic matrix. It can be used in the collection of HIV infection. The present sequence is a peptide of the V3 loop of HIV envelope protein gpl20. This peptide may be used in the composition of the present invention. gpl20 introducing the present eptide, the CD4 receptors are blocked, thereby introducing HIV infection. (Updated on 11-SEP-2003 to standardise os
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  Gaps
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    Indels
                                                                                                                                                                                                                                                                                                                                                  HIV gp120 V3 loop; liposome composition; HIV infection.
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Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                            HIV gp120 V3 loop peptide #1.
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                                      1 RIQRGPGRAFVTIGK 15
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(first entry)
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15; Conservative
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25-APR-2001
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Matches
  Matches
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AAU96031 standard; protein; 15 AA.
AAU96031
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RESULT 37

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Gaps

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The present invention relates to a new adjuvant of a gene derived from acid-fast bacterium for a DNA vaccine against AIDS (acquired immunodeficiency syndrome) and hepatitis C virus. The invention is composed of an effective component of alpha-antigen gene derived from acid-fast bacterium for DNA vaccine, particularly encoding for an alpha-antigen, particularly derived from Mycobacterium kanasali or its variant which has the same function, with an effective ingredient of an expression vector of the gene, used as an adjuvant, particularly a chimera DNA vaccine with a gene encoding for an antigenic peptide inserted, used for a DNA vaccine using a gene encoding for an immunogenic peptide derived from AIDS or hepatitis C virus. The adjuvant is useful for the treatment of AIDS or hepatitis C. The adjuvant enhances the immune inductive effect of the DNA vaccine. The present amino acid sequence represents the HIV (human immunodeficiency virus) CTL epitope peptide of the invention. (Updated on 29-AUG-2003 to standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       An adjuvant of a DNA vaccine composed of alpha-antigen genes derived from
                    Adjuvant; acid-fast bacterium; acquired immunodeficiency syndrome; DNA vaccinar, ALDS; hepatitis C virus; alpha-antigen gene; CTL; Mycobacterium kansasii; antigenic; immunogenic; epitope; human immunodeficiency virus; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eliciting an immune response; peptide antigen; T-cell epitope; tumour antigen; viral antigen; non-viral vector; HIV-1; T-cell co-stimulatory molecule; human immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 77; DB 5; Length 15; 100.0%; Pred. No. 9e-05; tive 0; Mismatches 0; Indela
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 7; 12pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABG68654 standard; peptide; 15 AA
                                                                                                                                                Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV-1 P18IIIB peptide antigen.
                                                                                                                                                                                                                                                                                           06-OCT-2000; 2000JP-00307674.
                                                                                                                                                                                                                                                                                                                                            06-OCT-2000; 2000JP-00307674.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                            (PRIM-) PRIMUNE CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acid-fast bacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunostimulant
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07-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    field)
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replicating vaccine delivery vehicle (which comprises a non-infectious
recombinant viral tubule) carrying one or more immunogens. The invention
is useful for inducing an immune response, preferably anti-viral, in a
cubject. The administration of the vaccine is preferably followed by
administering one or more virus like particles carrying an immunogen. It
is also useful for administrating to a patient for generating antibodies
specific for one or more immunogens, such as Bluetongue virus (BTV), foot
and mouth disease virus (FMDV), influenza virus and human fective means
confidency virus (HTV). The invention provides an effective means
of delivering multiple peptide components representing viral/tumour
antigenic groups to elicit protective immunity, which has not previously
contigenic groups to elicit protective immunity, which has not previously
contigenic groups to elicit protective immunity, which has not previously
contigent of the invention as immunogens. These epitopes were used to
construct chimeric NSI-TUBS (virus-derived tubules) which show
immunogenicity. (Updated on 29-NUG-2003 to standardise OS field)
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                                                                                                                                                                  Vaccine, non-replicating, viral tubule, immunogen, antibody, BTV;
Bluetongue virus, foot and mouth disease virus, FMDV, influenza virus,
human immunodeficiency virus, HIV; protective immunity, epitope, TUB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A vaccine, for inducing an antiviral immune response, comprises a non-replicating vaccine delivery vehicle (which comprises a non-infectious recombinant viral tubule) carrying one or more immunogens.
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                                                                                                                    HIV epitope, HIV-1 gp120 H-2Dd,help, peptide sequence.
                                                                                                                                                                                                                                              virus-derived tubule; anti-HIV; virucide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU97690 standard; peptide; 15 AA.
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                                                                                                                                                                                                                                                                                           Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                                         27-SEP-2001; 2001WO-US030464.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-SEP-2000; 2000US-0235614P.
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                                                (revised)
(first entry)
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(first entry)
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Matches 15; Conservative
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                                             29-AUG-2003
02-JUL-2002
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13-AUG-2002
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AAU96031;
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RESULT 39

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Roy P;

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Gaps

Berzofsky J;

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Claim 1; Col 78; 42pp; English
Khleif S,
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                                                                                                                                                                                                                           protein antigen comprising T-cell epitope(s) (e.g. tumour antigen, viral protein antigen comprising T-cell epitope(s) (e.g. tumour antigen, viral vortical antigen) coordinately with a non-viral vector comprising a polynucleotide encoding a T-cell co-stimulatory molecule. Viral peptide antigens may include human immunodeficiency virus (HIV) antigen, hepatitis B virus (HBV), herpes simplex virus (HSV) or human papilloma virus (HPV). The method is useful to elicit an immune response in a subject, and to supplement and enhance peptide and protein based vaccines and treatment methods. ABG68640-ABG68700 represent HIV-1 peptide antigens useful in the method of the present invention. (Updated on 29-AUG-2003 to
                                                                                                                                          Administering a non-viral vector encoding a co-stimulatory molecule alongside a peptide or protein T cell epitope, elicits increased response to the antigen and is useful to enhance peptide and protein based
                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                   present invention relates to a method for eliciting an immune
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eliciting an immune response; peptide antigen; T-cell epitope; tumour antigen; viral antigen; non-viral vector; HIV-1; T-cell co-stimulatory molecule; human immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 5; Length 15; 100.0%; Pred. No. 9e-05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus 1; (IIIB isolate)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV-1 P18 based peptide antigen #1.
                                                                                                                                                                                               Disclosure, Page 7; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABG68663 standard; peptide; 15 AA.
                       14-MAR-2001; 2001US-00810310.
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                                            15-MAR-2000; 2000US-0189396P
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                        15; Conservative
                                                                                                                                                                          vaccines and treatments.
                                                                                                Khleif S, Berzofsky J;
                                                                                                                                                                                                                                                                                                                                       standardise OS field)
                                                                 KHLEIF S.
BERZOFSKY J.
                                                                                                                    WPI; 2002-507231/54.
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(BERZ/) BERZOFSKY J.
                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
Matches 15; Conser
                                                                                                                                                                                                                                                                                                                                                             Sequence 15 AA;
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                                                                 (KHLE/) KHLEIF
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07-OCT-2002
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 18-APR-2002
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                                                                          (BERZ/)
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ABG68663
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                                                           Administering a non-viral vector encoding a co-stimulatory molecule alongside a peptide or protein T cell epitope, elicits increased response to the antigen and is useful to enhance peptide and protein based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant nucleic acid for the treatment and prophylaxis of acquired immunodeficiency syndrome, comprises a nucleic acid sequence encoding an antigen containing two or more cytolytic T lymphocyte (CTL) epitopes or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytolytic T lymphocyte; epitope; vaccine; prophylaxis; HIV infection; human immunodeficiency virus; acquired immune deficiency syndrome; CTL; gene therapy; AIDS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 77; DB 5; Length 15; 100.0%; Pred. No. 9e-05; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                              Disclosure; Page 7; 39pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus.
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                                                                                                                                                                   vaccines and treatments.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-148439/14.
WPI; 2002-507231/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV CTL epitope #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV; antigen; epitope; T cell; MHC; major histocompatibility complex; CTL; cytotoxic T lymphocyte; HIV infection; cancer; tuberculosis; tumour; hepatitis; melanoma; breast cancer; Hodgkin lymphoma; nasopharyngeal carcinoma; vaccine; immune response; hyaluronic acid; HA; CD8+ T cell; CD8+ T cell; viral infection; bacterial infection; tungal infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modulating an immune system response to an antigen in a mammal, comprises administering a particle-free therapeutic comprising a hyaluronic acid polymer analogue covalently linked to a peptide that comprises a T cell
           The invention relates to a recombinant nucleic acid comprising a nucleic acid sequence encoding an antigen containing two or more cytolytic T lymphocyte (CTL) epitopes or its analoques. Sequences of the invention are used in vaccines and are useful for the treatment and prophylaxis of human immunodeficiency virus (HIV) infection, particularly acquired inmuno deficiency syndrome (AIDS). The invention is also useful in gene therapy. The present sequence is HIV CTL epitope. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                      100.0%; Score 77; DB 6; Length 15; 100.0%; Pred. No. 9e-05; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ADN14074 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV helper T cell epitope #41.
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                                                                                                                                                                                                                                                                                                   1 RIQRGPGRAFVTIGK 15
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Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chu Y, Qiu J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MPI; 2003-540464/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JS2003049253-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LI F O.
                                                                                                                                                                                    Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oru J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADN14074;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHUY/)
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ID ADN
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the mammal, or an MHC Class II molecule and by a CD8+ T cell of the mammal, or an MHC Class II molecule and a CD4+ T cell of the mammal.

C The immune system response comprises a cytotoxic T lymphocyte, a CD4+T cell, or an antibody that recognises the antigen. The immune system response to the antigen is increased after administration of the antigen is increased after administration of the antigen is an antigen of a pathogenic agent or a tumour cell. The immune system response to the antigen is an antigen of a tissue or organ transplanted to the mammal. The composition and methods care useful for modulating, i.e. enhancing or diminishing, an immune system response to an antigen in a mammal. The composition is also useful configurates are useful as peptide vaccines against an antigen, a conjugates are useful as peptide vaccines against an antigen, a conjugates are useful as peptide vaccines against an antigen, a conjugates are useful as peptide vaccines against an antigen, a cuseful for treating or preventing diseases associated with any of the antigens above e.g. HIV infection, cancer, tuberculosis, hepatitis, antigens above e.g. HIV infection, a lymphoma and nasopharyngeal carcinoma. The peptide vaccine compositions are additional adjuvants, but still induce a stronger cell-mediated response than peptide vaccines of the present sequence is an HIV-1 cell of the prior art. The present sequence is an HIV-1 cell or the vaccine of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New vaccine composition comprising a peptide adjuvant and a DNA vaccine encoding an immunogenic protein, useful for inducing immune responses against diseases e.g. HIV infection, autoimmune diseases, tuberculosis or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kim Y, Sung Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vaccine; virucide; antibacterial; immunosuppressive; antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 77; DB 7; Length 15; 100.0%; Pred. No. 9e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immune response induction composition peptide adjuvant #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lee C, Choi S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYPO-) UNIV POHANG SCI & TECHNOLOGY.
(GENE-) GENEXINE CO LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 21; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADR04041 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-JAN-2004; 2004WO-KR000177.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2003; 2003KR-00006393.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; peptide adjuvant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Park K, Park S, Yang S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-AUG-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    allergies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADR04041;
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The present invention relates to a vaccine composition comprising a

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2 RIORGPGRAFVTIGK 16

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             composition may also comprise a gene of the influence virus, preferably the nouraminidase gene. The vaccine composition is useful for inducing immune responses against diseases comprising HIV infection, herpes simplex virus (HSV) infection, influence virus infection, hepetitis and a infection, papillomavirus infection, tuberculosis, tumour growth, autoimmune diseases or allergies. The present sequence is a peptide adjuvant useful in the composition of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lipopeptide, lipoprotein; vaccine; cytotoxic T-cell; lymphocyte; HIV; human immunodeficiency virus; AIDS; cancer; tumour cells; CB1; CB2; CB3.
peptide adjuvant and a DNA vaccine encoding an immunogenic protein. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The sequence is that of peptide ENV 312-327 derived from the HIV, it is
                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lipopeptide(s) which stimulate cytotoxic T-cells - for treating HIV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gras-Masse H;
                                                                                                                                                                                                                  ;
                                                                                                                                                                              100.0%; Score 77; DB 8; Length 15; 100.0%; Pred. No. 9e-05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sergheraert C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (INSP ) INST PASTEUR LILLE.
(INRM ) INSERM INST NAT SANTE & RECH MED.
                                                                                                                                                                                                                                                                                                                                                                 AAR24939 standard; protein; 16 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  parasitic infections and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 18; 32pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Levy JP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               91EP-00403446.
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                                                                                                                                                                                                                                                 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
                                                                                                                                                                             Query Match
Best Local Similarity 100.0
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Martinon F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV peptide ENV 312-327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) INST PASTEUR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Boutillon C, Marcan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-209776/26.
                                                                                                                                                Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
09-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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Peptides AAW68301-W68384 are examples of antigens (Ag) which can be loaded onto recombinantly produced major histocompatibility complex (MHC) molecules in a method of detecting antigen-specific lymphocytes. The MHC-antigen complex is then immobilised on a solid support and a sample containing cells recognising the MHC-Ag complex may be isolated. This peptide is derived from amino acids 312-327 of the human immunodeficiency virus type 1 (HIV-1) env protein. A similar method is used to isolate, purify or eliminate Ag-specific T-cells or to produce Ag-specific cytotoxic T-cells (CTC). The method is also used to detect and quantify tumour-specific T-cells and to generate CTC for specific killing of tumour specific (solid tumours, leukaemia or lymphoma) by injection into a human or animal, but also for treating viral infections. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detection, purification and elimination of antigen-specific lymphocytes for producing cytotoxic T cells for immuno-therapy of cancers and viral infection.
                                                                                                                                                                               complex; MHC; lymphocyte; detection;
tumour; leukaemia; lymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 2; Length 16; 100.0%; Pred. No. 9.5e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Langladedemoyen P, Lone Y, Kourilsky P, Abastado J;
                                                                                                                                                                                                                                                                                                                                                                                                                             (INSP ) INST PASTEUR.
(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 27; 222pp; French.
                               AAW68326 standard; peptide; 16 AA.
                                                                                                                                                                              Antigen; major histocompatibility immobilisation; cytotoxic T-cell;
                                                                                                                                                MHC binding peptide Env.312-327.
                                                                                                                                                                                                                                                                Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                 97WO-FR000892.
                                                                                                                                                                                                                                                                                                                                                                                                96US-00651925.
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                                                                                             (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-018653/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                viral infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16 AA;
                                                                                                                                                                                                                                                                                                                                                               21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                21-MAY-1996;
                                                                                                                                                                                                                                                                                               WO9744667-A2
                                                                                              25-MAR-2003
14-OCT-1998
                                                                                                                                                                                                                                                                                                                                27-NOV-1997
                                                                                                                                                                                                                                               Synthetic.
                                                               AAW68326;
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Matches
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AAY68203
ID AAY68
RESULT 46
               AAW68326
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AAY68203 standard; peptide; 16 AA.

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Gaps

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Length 16; Indels

Query Match
100.0%; Score 77; DB 2; L
Best Local Similarity 100.0%; Pred. No. 9.5e-05;
Matches 15; Conservative 0; Mismatches 0;

us-08-869-386-1.rag

Altered MHC determinant binding peptide SEQ ID NO:35.

(first entry)

14-FEB-2000

AAY52857;

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New recombinant DNA encoding covalently linked form of major histocompatibility complex Class I determinant, used for immune system stimulation, e.g. for treating cancer.
                                                                       MHC class I; major histocompatibility complex; microglobulin; antigen; immune response; immunisation; AIDS; multiple sclerosis; toxic shock; cancer; lupus erythematosus; snake bite; cytostatic; antiviral; immunomodulatory; dermatological; immunosuppressive; antinflammatory;
                                                        Altered MHC determinant binding peptide SEQ ID NO:35.
                                                                                                                                                                                                                                         (INSP ) INST PASTEUR.
(INRM ) INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                                                        Abastado J;
                                                                                                                                                                                                                                                                                                                                                    Disclosure, Col 11, 88pp; English.
                                                                                                                                   Human immunodeficiency virus 1.
                                                                                                                                                                                            95US-00481985
                                                                                                                                                                                                               91US-00792473
91US-00801818
                                     (first entry)
                                                                                                                                                                                                                                                                         Kourilsky P, Mottez E,
                            (revised)
                                                                                                                                                                                                                                                                                           WPI; 2000-125951/11.
                                                                                                                  neuroprotective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 16 AA;
                                                                                                                                                                                                               15-NOV-1991;
05-DEC-1991;
                                                                                                                                                                                             07-JUN-1995;
                           12-SEP-2003
                                     13-APR-2000
                                                                                                                                                      US6011146-A
                                                                                                                                                                          04-JAN-2000
         AAY68203;
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The present invention describes a recombinant DNA molecule (I) containing a sequence (Ia) that encodes an altered MHC (major histocompatibility complex) Class I determinant (II) comprises a polypeptide with alphal, alpha2, alpha3 and beta2-microglobulin domains, in which alpha3 and beta2 are covalently linked, thorough C- and N-termini respectively, via a nucleotide spacer sequence encoding a polypeptide. (II) includes an artigen-binding site and when (II) and the antigen are associated they are recognized by a mammalian T cell receptor (TCR). (I) are used to produce (II) which are used to study functional interactions between the various MHC domains. They can also be used to modulate (in vivo or in vitro) the immune system by inducing an effector response (cytotoxicity, antibody synthesis, phagocytosis) of immune system cells, typically for treating, or immunising against; cancer, acquired immune deficiency syndrome, lupus erychematosus, multiple sclerosis, toxic shock and snake bite, but also for selective destruction of autoreactive cells, diagnosis, therapy, studying MHC-associated cellular processes and for affinity purification). AASE3758 and AAY68186 to AAY6816 are sequences used in the exemplification of the present invention. (Updated on 12-SEP-2003 to standardise OS field)
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                               Gaps
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100.0%; Score 77; DB 3; Length 16; 100.0%; Pred. No. 9.5e-05; ive 0; Mismatches 0; Indels
                                                             RIORGPGRAFVTIGK 15
               Local Similarity 100.
108 15; Conservative
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   Query Match
                               Matches
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RIORGPGRAFVTIGK 16
                                  RESULT 48
                                             AAY52857
ID AAY
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AAY52857 standard; peptide; 16 AA.

AAB58618 standard; peptide; 16 AA.

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The present invention describes a composition capable of eliciting antimajor histocompatibility (MHC) antibodies. The composition comprises an antigen associated with an altered MHC class II determinant (I) comprising alphal, alpha2, betal and beta2 polypeptide domains encoded by a mammalian MHC class II locus covalently linked to form a polypeptide (I) containing beta2, alpha2, alpha1 and beta1 domains in sequence. The resulting Antigen-MHC complex is recognizable by the T cell receptor: The compositions are used for immunisation against, or treatment of, a wide range of autoimmune diseases, e.g. acquired immune deficiency syndrome (ILDS), lupus erythematosus, multiple sclerosis, thyroiditis, toxic shock, tumour and snakebite, depending on the nature of antigen. (I) is also used to analyse functional interactions between the various domains and for targeting lymphocyte receptors. Antibodies against (I) are produced by usual methods of immunisation or cell fusion, and may be humanised by standard methods. These antibodies are useful for diagnosis (detection or purification of MHC gene products), therapy (neutralishing MHC on cell surfaces) and in the study of MHC and cellular processes:

AAZ33240 to AAZ33242 and AAX52840 to AAX52970 represent sequences used in the exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Composition containing an antigen and altered major histocompatibility Class II determinant, used to immunize against autoimmune diseases, e.g.
                                                                                                                             Major histocompatibility complex; MHC class I; MHC class II; antigen; immune response; disgnosis; antibody; immunisation; autoimmune disease; acquired immune deficiency syndrome; AIDS; cytostatic; dermacological; anti-inflammatory; neuroprotective; immunosuppressive; antithyroid; vaccine; lupus erythematosus; multiple sclerosis; thyroiditis; toxic shock; tumour; snakebite.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (INRM ) INSERM INST NAT SANTE & RECH MEDICALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Class II determinant, used to immuni:
acquired immune deficiency syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kourilsky P, Mottez E, Abastado J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Col 11; 96pp; English.
                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                        95US-00484905.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91US-00801818
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUN-1995;
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                                                                                                                                                                                                                                                                                  Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AABS8618
ID AARS
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AAB58618;

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Peptide P-18, opt. with a Cyg-Gly dipeptide attached at its N- terminus, is the pref. peptide for use in raising a cytotoxic T lymphocyte response against HIV. The peptide is encapsulated in a liposome for admin. (Updated on 2-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Inducing cytotoxic T lymphocyte response to HIV - with liposome contg. peptide or protein having CTL epitope of HIV and protein, also improving numeral immunity, useful in vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "this residue is -NH-CHR-CO-NH2, where R is a Cl4
                                        immunogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lipopeptide, antigen; cytotoxic T lymphocyte; steroid; vaccine; HIV related condition; tumor cell; NP protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                /note= "the Cys-Gly dipeptide is opt. absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 77; DB 2; Length 17; 100.0%; Pred. No. 0.0001; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lipopeptide comprising a fragment of the HIV env protein.
                                   Human Immunodeficiency Virus type 1; envelope protein; vaccine; AIDS; peptide P18; epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                3
Peptide CG-P18 from HIV-1 IIIB env protein V3 loop
                                                                                                                                                                                                                                                                                                                                                                                                                Wassef N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "amidated residue"
                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                Koenig S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY40414 standard; peptide; 17 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 16; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                         92US-00860707.
                                                                                                                                                3. .17
/label= P-18
                                                                                                                                                                                                                                                                                      93WO-US002978.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 RIORGPGRAFVTIGK 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              side chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                Cassatt D,
                                                                                                                                                                                                                                                                                                                                                            MEDI-) MEDIMMUNE INC
                                                                                                                                                                                                                                                                                                                                                                             (USSA ) US DEPT ARMY
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1993-336590/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modified-site
                                                                                                                                                                                                                                                                                      25-MAR-1993;
                                                                                                                                                                                                                                                                                                                         31-MAR-1992;
                                                                                                                                                                                                               W09319775-A1
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                                                                                                                                                                                                                                                  14-0CT-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                Alving CR,
                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY40414;
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Best Local S
                                                                                                                                            Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a major histocompatibility complex (MHC) class I determinant, which has alpha 1 alpha 2 alpha 3 and beta2-
microglobulin polypeptide domains encoded by a mammalian MHC class I locus. The MHC class I determinants ere useful for activating the immune system and presenting antigens to the immune system to elicit an antigenic response. The MHC class I determinants are also useful for treating diseases, e.g. T cell mediated autoimmune disease, AIDS, lupus erythematosus, toxic shock or snake bite. The altered MHC class I determinants and compositions containing antigens bound to the determinants may be useful in diagnostic applications, e.g. altered determinant can be assayed. (Updated on 11-SEP-2003 to standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New altered major histocompatibility complex (WHC) class I determinant useful for eliciting an immune response and/or for immunizing against or treating diseases, for example, multiple sclerosis, AIDS, toxic shock or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mottez E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                       Major histocompatibility complex; MHC class I; immune; snake bite; T cell mediated autoimmune disease; AIDS; lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ojcius D, Lone Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 77; DB 4; Length 16; 100.0%; Pred. No. 9.5e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                         Altered MHC determinant binding peptide #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Casrouge A,
                                                                                                                                                                                                                                                                                                                                                                                                                                              (INSP ) INST PASTEUR.
(INRM ) INST NAT SANTE & RECH MEDICAL.
                                                                                                                                                                                                             Human immunodeficiency virus; type 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Col 11; 105pp; English.
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91US-00801818.
93US-00072787.
93US-00117575.
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                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Abastado J, Kourilsky P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
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                                                     (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-060089/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 16 AA;
                                                                                                                                                                                                                                                                                                                                                                       05-DEC-1991;
07-JUN-1993;
07-SEP-1993;
                                                                                                                                                                                                                                                                                                                       09-JAN-1995;
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29-APR-1994
                                                  11-SEP-2003
13-MAR-2001
                                                                                                                                                                                toxic shock.
                                                                                                                                                                                                                                                  US6153408-A.
                                                                                                                                                                                                                                                                                                                                                          15-NOV-1991;
                                                                                                                                                                                                                                                                                      28-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  snake bite.
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AAR42057;

RESULT 50
AAR42057
ID AAR4
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AC AAR4
XX
DT 25-M
XX

field

Query Match Matches

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Gaps

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Smini A, Liu MA, Marburg S,
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                                                                                              WPI; 1992-026505/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
28-APR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP519554-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR30032;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR30032
엄
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "bonds to the OMPC of the conjugate via this site"
                                                                                                                                                                                                                                                                                                                                                                                                                The specification describes lipopeptide that comprise a partial peptide containing 10-40 amino acids and at least one antigenic determinant specific for cytotoxic Tlymphocytes. The lipopeptide comprises at least one 10-20 carbon fatty acid derivatives and/or at least one modified steroid group. The lipopeptides are useful for: the preparation of a vaccine against HIV related conditions; immunitaing a human or animal against tumor cells; and immunizing human or animal against tumor cells; and immunizing human or animal against tumor cells; and immunizing human or animal parasaites). The present sequence represents a lipopeptide of the invention, and comprises part of the HIV env protein
                                                                                                                                                                                                                                                                                                                  New lipopeptide inducers of cytotoxic {\tt T} lymphocytes, useful as vaccine against cancers, viruses, parasites and HIV-related conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                   Gras-Masse H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AIDS; ARC; human immunodeficiency virus; vaccine; Neisseria; meningitidis b; outer membrane protein complex; OMPC; PND135-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 77; DB 2; Length 17; 100.0%; Pred. No. 0.0001; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                 Boutillon C, Martinon F, Sergheraert C, Magne R,
Gomard E, Tartar A, Levy J;
                                                                                                                                                       (INSP ) INST PASTEUR LILLE.
(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV principal determinant peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR31277 standard; peptide; 18 AA
                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 19; 35pp; French
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90US-00555966.
91US-00715276.
91US-00715278.
                                                                                              90FR-00015870.
                                                        99EP-00105773
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 RIORGPGRAFVTIGK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-FEB-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 100.
nes 15, Conservative
                                                                                                                                                                                                                                                                              WPI; 1999-553128/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 AA;
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Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUL-1990;
19-JUL-1990;
19-JUN-1991;
19-JUN-1991;
                                                        18-DEC-1991;
                                                                                                18-DEC-1990;
18-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-JUL-1991;
                  29-SEP-1999
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Matches
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IID AAR3

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The invention relates to a co-conjugate comprising an immunogenic protein or protein complex having a first set of covalent linkages to low molecular weight moieties which have an anionic or polyanionic character at physiological pH, and a second set of covalent linkages to peptides comprising HIV principal neutralizing determinants (PND's) or immunologically equivalent peptides. Preferably at least one set of the covalent linkages is comprised of maleimide derivatives; the covalent linkages is composed of one to five residues of the anionic covalent linkages is composed of one to five residues of the anionic covalent linkages is composed of one to five residues of the anionic covalent linkages is composed of one to five residues of the anionic covalent linkages is composed of one to five residues of the anionic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b; and the PND peptide has a linear structure, a disulphide-bonded cyclic structure or a thiosether-bonded cyclic structure. The present sequence (PND135-18) is an example of a PND peptide component used in the co-conjugate. The co-conjugate is useful for inducing anti-peptide immune response in mammals, for inducing HIV infection or disease, including AIDS, or for treating humans afflicted with HIV infection or disease.
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major immune enhancing protein; vaccine; anti-HIV antibodies; immunogen;
passive immunisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                        Conjugate proteins comprising HIV peptide components - useful for preparing vaccines for e.g. AIDS or for treating infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV principle neutralising determinant 135-18.
                                                                                       Emini EA;
                                                                                       Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR30032 standard; peptide; 18 AA.
                                                                                                                                                                                                                                                                                                                                                                                         Claim 12; Page 56; 63pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15; Conservative
                                                                                       Leanza WJ, Marburg S,
(MERI ) MERCK & CO INC.
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Tolman RL;

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The peptide is HIV principle neutralising determinant (PND) 135-18 and is used as part of a conjugate comprising the major immune enhancing protein (MIEP) of Neisseria meningitidis covalently linked to the HIV PND. The conjugate may be used to prepare vaccines against HIV infections, e.g. AlbS, as research tools for studying PND structure-function relationships, or as immunogens for use in the passive immunisation of humans. (Updated on 25-MAR-2003 to correct PN field.)
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C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "Joins onto polysaccharide-protein complex via
this site"
Conjugates of HIV-1 PND peptide(s) with the MIEP of Neisseria meningitidis - useful as a vaccine for treating and preventing HIV-1 infection, e.g. AIDS in humans.
                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV peptide-polysaccharide-protein conjugates - used in vaccines or produce antibodies to prevent or treat HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus; principal neutralizing determinant;
outer membrane protein complex; OMPC; Neisseria; AIDS; PND-135-18.
                                                                                                                                                                                                                           ö
                                                                                                                                                                                                  Length 18;
                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                Score 77; DB 2; L
Pred. No. 0.00011;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                         HIV-PND-polysaccharide-protein conjugate vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EA;
                                                Claim 9; Page 59; 66pp; English.
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                                                                                                                                                                                                                                                                                                                                  AAR26713 standard; peptide; 18
                                                                                                                                                                                                 100.0%;
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90US-00555974.
91US-00715275.
91US-00715277.
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                                                                                                                                                                                                                                               1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                              1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                        15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1992-034437/05.
                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                          Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUL-1990,
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                                                                                                                                                                                                                                                                                                                                                         AAR26713;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                  Query Match
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The invention relates to a conjugate of an HIV principal neutralizing determinant (PND), or an immunologically equivalent peptide (PEP), covalently coupled to an immunogenic protein or protein complex through an anionic polysaccharide linker. Pref. the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b and the PND peptide has a linear structure, a disulphide-bonded cyclic structure,

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an amide-bonded cyclic structure or a thioether-bonded cyclic structure. The present sequence (PNU13-18) is an example of a PND peptide component. The conjugates are used for inducing HIV-neutralising antibodies or for making vaccines to prevent contraction of HIV infection of disease. The antibodies can be used for passively protecting against infection by HIV, or for protecting against proliferation of HIV postinfection, or for treating AIDS, or in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequence given in AAR44190 is a peptide antigen, B2, which represents residues 112-329 of the third variable domain (V3 loop) of gp120, of HIV-18 train IIIB. This sequence was attached to an amino acid linker (see also AAR44191) in the production of a multiple antigenic peptide system. This system comprises a dendritic core to which are covalently attached at least one peptide, eg. an antigenic peptide, and a lipophilic membrane anchoring group. This system may be injected into a mammal and elicits both humoral and cytotoxic T cell (CTL) immune responses. This system may be used to immunise against HIV infection. The lipophilic membrane anchoring group provides efficient adjuvant activity without the toxicity purblems of Freunds adjuvant, while the dendritic structure allows multiple antigens to be attached. Optionally the antigens may be derived from different pathogens, providing vaccines which protect against more than one disease, eg. HIV, influenza and malaria. (Updated on 25-MAR-2003
                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antigen, B2; third variable domain, V3 loop; gp120; HIV-1; vaccine; strain IIIB; multiple antigenic peptide system; dendritic core; lipophilic membrane anchoring group; mammal; humoral; immunisation; cytoctoxic T cell; CT; immune response; infection; Freunds adjuvant; pathogen; HIV; influenza; malaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New multiple antigen system esp. for use in HIV vaccines - contains lipophilic membrane anchor imparting adjuvant activity, and peptide antigens coupled to dendritic core.
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                                                                                                                                                                                                     Length 18;
                                                                                                                                                                              100.0%; Score 77; DB 2; Length l8
100.0%; Pred. No. 0.00011; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   AAR44190 standard; peptide; 18 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                                                                                    1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                               Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                             1 RIQRGPGRAFVTIGK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gp120 V3 loop antigen B2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1993-368723/46.
                                                                                                                                                            Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-MAY-1993;
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20-MAY-1994
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AAR58548

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Vaccine for immunization against tumor cells or pathogens, especially HIV, comprising peptide part, antigenic determinant specifically inducing cytotoxic T-lymphocytes and N-palmitoyl-lysine-derived chain(s).
                                                                                                                                                                                                                             /note= "xaa is optionally 2-acetylamino-hexadecanoyl, 2,4-bis(hexadecanoylamino)butyryl, or not present"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is a lipopeptide, which can be used for the immunisation of humans or animals against tumour cells or pathogens, specifically viruses, especially HIV-1 or HIV-2. The pathogens may also include parasites. Examples illustrate immunisation of mice against influenza, as well as HIV. The lipopeptide, with the appropriate response in a host organism against a strong cytotoxic T-lymphocyte response in a host organism against a wide range of pathogens. Addition of an adjuvant is unnecessary
                                                                                            Lipopeptide; cytostatic; virucide; anti-HIV; antiparasitic; vaccine; immunisation; tumour; pathogen; virus; antiviral.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gras-Masse H;
                                                                                                                                                                                                                                                                                             /note= "Xaa is optionally 2-amino-hexadecanoamide epsilon-hexadecanoy1-Lys"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               a F, Sergheraert C, Magne R,
Levy J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 77; DB 4; I 100.0%; Pred. No. 0.00011;
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                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR60203 standard; protein; 20 AA.
                                                             Lipopeptide #2 used in a vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 4; Page 17; 31pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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99EP-00105773.
                                                                                                                                                                                                                                                                                                                                                                                                                  18-DEC-1991; 2000EP-00117513
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                                                                                                                                                                                                                                                                                 /label= Xaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INSP ) INST PASTEUR LILLE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Martinon F,
                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Boutillon C, Martino
Gomard B, Tartar A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-114040/13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 AA;
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Wodified-site
                                                                                                                                                                                                                                                                 Modified-site
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18-DEC-1991;
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                              05-AUG-2002
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25-MAR-2003
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                                                                                                                                                Synthetic
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 ABB83113;
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Matches
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ID AAR6
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AC AAR6
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DT 27-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (AARSB550) of HIV env protein, the domain being capable of inducing a helper T cell response, and 5 peptide domains from the V3 loop of HIV-1 isolates MN, SC, RF, IIIB and WNJ2 (AARSB45-49), each peptide being separated by a spacer domain (AARSB551). The gene was expressed in E. CALLINKK-MRNY for preparation of a multiple epicope AIDS vaccine (AARSB552). (Updated on 25-WAR-2003 to correct PN field.) (Updated on 16-OCT-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1; V3 loop, multiple epitope; AIDS; vaccine; MEAV; Escherichia coli;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New recombinant proteins contg multiple antigenic determinants - linked by flexible hinge domains.
to correct PN field.) (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene (AAQ70535) encodes a portion of the CD4 binding domain
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                                                                 Length 18
                                                                                              0; Indels
                                                                 Score 77; DB 2; I Pred. No. 0.00011;
                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 36; 56pp; English.
                                                                                                                                                                                                                                                                                                                                                                               HIV-1 isolate IIIB V3 loop domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABB83113 standard; peptide; 18 AA.
                                                                                                                                                                                                                                                    AAR58548 standard; peptide; 18 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1
                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UNBI-) UNITED BIOMEDICAL INC.
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                                                                                                                                                               4 RIORGPGRAFVTICK 18
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                                                                               Local Similarity 100.
nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                  (revised)
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Matches 15; Conserv
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                                Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-FEB-1993;
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29-MAR-1995
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                                                                 Query Match
                                                                                                  Matches
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Gaps

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ABB83113

88US-00155321

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Example 1; SEQ ID NO 37; 127pp; English.
          01-MAR-1991;
09-JUL-1991;
                              19-OCT-1994;
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                                                                                                                  components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                                      ADR18886;
                                                                   Wang CY;
                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                              A molecular tag was created by annealing two complementary 76mer oligonucleotides with cohesive end overhangs. AA070167 is the sense strand and includes a Bcl coverhang, the HIV gpl10 v3 loop coding sequence and a stop codon. The peptide encoded by the molecular tag sequence and a stop codon. The peptide encoded by the molecular tag regions from different antibodies, affected the avidity and binding specificity of the antibodies. For example, the tag failed to function properly when fused to L6 but performed successfully when fused to CD3. (Updated on 25-MMR-2003 to correct OS field.)
                                                                                                                                                                                                                                       Expression vector encoding bispecific fusion protein - having binding domains for separate targets joined by helical peptide, useful e.g. for diagnosis and treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunoadsorbent; immunoassay; HIV gp120; immunogen; antibody; Human.
                                                                                                                                                                                           Bajorath J;
                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                      fusion protein; recombinant bispecific single chain antibody; human immunodeficiency virus; glycoprotein gp110; V3 loop.
                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 77; DB 2; Length 20; 100.0%; Pred. No. 0.00012; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                          Linsley PS,
                                                                                                                                                                                         Hayden MS,
                                                                                                                                                                                                                                                                             Example 1; Page 12; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW54930 standard; peptide; 20 AA.
                   HIV gp110 V3 loop molecular tag.
                                                                                                                                                                       (BRIM ) BRISTOL-MYERS SQUIBB CO.
                                                                                                                                          93US-00013420.
93US-00121054.
                                                                                                                         94EP-00300692.
                                                                                                                                                                                         Ledbetter JA, Gilliland LK,
                                                                 Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19
13-MAR-1995 (first entry)
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Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 AA;
                                                                                                                        31-JAN-1994;
                                                                                                                                          01-FEB-1993;
13-SEP-1993;
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                                                                                                      10-AUG-1994.
                                                                                   EP610046-A2
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Н
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptides AAW54903-W54941 can be used as an immunoadsorbent in an immunoassay for detecting antibodies to HIV gpl20, or as an immunogen for eliciting antibodies to HIV in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Composition for inhibiting HIV-1 infection, comprises isolated peptide molecule that mimics atomic structural conformation of V3 loop peptide of HIV-1 envelope glycoprotein that is bound to, and constrained by human monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     three-dimensional atomic structural conformation; protein co-ordinate data; V3 loop peptide; HIV-1; envelope glycoprotein; gp120; human monoclonal antibody 447-52D; murine monoclonal antibody 0.5 beta; immunogen; immunogenic; V3 loop epitope; HIV-1 infectivity inhibitor; anti-HIV; vaccine; HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                     HIV gp120 peptides - useful as immunoassay reagents or vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 77; DB 2; Length 20; 100.0%; Pred. No. 0.00012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anglister J, Sharon M, Schapira M, Zolla-Pazner S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV-1 V3-IIIB related peptide SEQ ID NO:37.
                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Column 21/22; 34pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADR18886 standard; peptide; 20 AA.
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                                                                                                                      (UNBI-) UNITED BIOMEDICAL INC.
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(YEDA ) YEDA RES & DEV CO LID.
91US-00663262.
                                                           94US-00326676.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 100.
nes 15; Conservative
                                                                                                                                                                                                                                         WPI; 1998-347301/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 AA;
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94US-00290105.

15-AUG-1994;

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The present invention describes a composition (CI) which comprises an isolated peptide molecule or isostere that minics the three-dimensional collected peptide molecule or isostere that minics the three-dimensional convenience of structural conformation of the VI) loop peptide of the HIV-I convenience by and constrained by, human monoclonal antibody (MAD) 447-52b, murine MAD to 5 beta or an antigen binding fragment of the MAD, where the constrained VI loop peptide of a molecule that is useful as an HIV-I VI loop peptide when it is in free form. Also described: (I) identifying (MI) from several existing compounds a molecule that is useful as an HIV-I VI loop immunogen or as an inhibitor of binding of HIV-I to a chemokine receptor/HIV-I correceptor on the surface of a receptor-bearing target cell; (2) designing a molecule that is useful as an HIV-I VI loop immunogen or as an inhibitor of binding of HIV-I to a chemokine receptor/HIV-I correceptor on the surface of a receptor-bearing target cell; (3) a composition (C2) that is useful as an HIV-I VI loop immunogen or as an inhibitor of binding of HIV-I to a chemokine receptor/HIV-I correceptor on the surface of a receptor-bearing target cell; (4) an immunogenic composition (C3) for induction of an anti-HIV-I antibody response specific for a VI loop peptide for a comprising (C2) and an actipient; (5) a paramaceutical composition (G3) and a carrier or excipient; (6) an immunogenic composition (C3) and a carrier or excipient; (6) a computing platform for generating a produce or excipient; (6) a computing platform for generating and peptide that is bound to 447-520 or 0.5beta MAD or 1.5beta MAD or 1.5b
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                           Gaps
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  Length 20;
                        0; Indels
100.0%; Score 77; DB 8; L
100.0%; Pred. No. 0.00012;
ive 0; Mismatches 0;
                                                   1 RIORGPGRAFVTIGK 15
                           Conservative
            Local Similarity
nes 15; Conserv
   Query Match
                         Matches
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Antigen; non-infectious; retrovirus; antigenic marker; immune response;
long terminal repeat; gag; pol; env; AIDS; HIV; antibody; therapy.
                                                   AAR93073 standard; peptide; 21 AA.
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   6 RIORGPGRAFVTIGK 20
                                                                                         27-SEP-1996 (first entry)
                                                                                                           Antigenic peptide CLTB73.
                                                                                                                                                                                                                       15-AUG-1995;
                                                                                                                                                                               WO9605292-A1
                                                                                                                                                                                                   22-FEB-1996
                                                                                                                                                            Synthetic
                                                                      AAR93073;
                                RESULT 61
                                         AAR93073
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           curative) treatment of AIDS and related conditions, by acting to displace the binding of the HIV virus to human or animal cells, or by disrupting the 3-dimensional organisation of the virus. The particle can also be used to identify antibodies specifically reacting with retrovirus
                                                                                                                               Antigenically marked non-infectious retrovirus-like particles - used to vaccinate against, and in the treatment of, AIDS and AIDS related
                                                                                                                                                                                                                                                                                                                                                                                                      preferably oral, anal, vapalal or intranasal administration. The composition can be used for immunising a host to produce a retroviral specific immune response, such as against AIDS and AIDS related conditions. The particles may also be used in the prophylactic (or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polypeptide N-acetylgalactosaminyltransferase; GalNAc-t3; human; glycosylation; HIV-V3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 2; Length 21; 100.0%; Pred. No. 0.00012;
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                                                                 Persson R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW34475 standard; peptide; 21 AA.
                                                                                                                                                                                                    Example 4; Page 38; 75pp; English.
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                                                                 Yao F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIORGPGRAFVTIGK 15
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                                 (CONN-) CONNAUGHT LAB LTD
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Best Local Similarity 100.*
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Acceptor peptide HIV-V3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CLAU/) CLAUSEN H.
(BENN/) BENNETT E P.
                                                                   Cao S,
                                                                                                 WPI; 1996-139690/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 AA;
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                                                                   Rovinski B,
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                                                                                                                                                                     conditions.
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                                                                                                                                                                                                                                    host-blood group O2 gene. Gallmac-transferase activity with the Muc2 acceptor peptide was increased 20-fold, and activity with the HIV-V3 peptide was increased nearly 100-fold. In contrast, expression of GalNac-T1 and -T2 constructs only increased the GalNAc-transferase activity toward Muc2 and Muc5C peptide substrates. This illustrates the unique acceptor substrate specificity of GalNAc-T3. The enzyme is used in claimed methods for the glycosylation of peptides and proteins and for producing vaccines by modifying the 0-glycosylation pattern of eukaryotic
                                                                                                                                               Acceptor peptides Muc2, Muc5c (see AAW34474) and HIV-V3 (see AAW34475) were used to study the acceptor substrate specificity of the novel human N-acetylgalactosaminyltransferase GalNAc-T3 (see AAW34470). Expression of a soluble GalNAc-T3 construct in Significally resulted in significant increases in GalNAc-transferase activity in the culture medium of infected cells compared to uninfected controls or cells infected with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   V3 loop; gp120 protein; HIV-1; retrovirus-like particle; genome; HIV-2; long terminal repeat; LTR; chimeric; envelope; glycoprotein; HTLV-1; HTLV-II; vaccine; human T-lymphotropic virus.
                                                         New isolated N-acetyl-galactosaminyl-transferase enzyme - used for the production of glycosylated polypeptide(s) having particular enzymatic, immunogenic or other biological or physical properties.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunogenic retrovirus-like particle - with chimeric HIV-1 envelope protein containing heterologous retroviral amino acid sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 77; DB 2; Length 21; 100.0%; Pred. No. 0.00012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
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                                                                                                                   Example 2; Page 30; 70pp; English
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93US-00073526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RIQRGPGRAFVTIGK 15
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(revised)
(first entry)
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 Bennett EP
                               WPI; 1998-008874/01
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ses 15; Conser
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 AA;
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09-JUN-1993;
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20-MAR-2003
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 Clausen H,
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Matches
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                                       This sequence represents a peptide from the V3 loop of the gpl20 protein from the human immunodeficiency virus type 1 (HIV-1) strain HXB2. The peptide is used to determine antibody responses after immunisation with a self-assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle. The retrovirus-like particle comprises a modified HIV genome devoid of long terminal repeats (LTRS) and contains a nucleotide sequence coding for a chimeric envelope glycoprotein. The chimeric envelope glycoprotein has the HIV-1 gpl20 conserved region 2 and a second retroviral envelope amino acid sequence from a heterologous strain of HIV-1, HIV-2, HILV-1 inserted into the first retroviral envelope amino acid sequence from a heterologous strain of HIV amino acid sequence (see AAW75474-W7547). The novel retrovirus-like particle is useful in vaccines against HIV. (Updated on 20-MAR-2003 to correct PA field.) (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes a nucleic acid molecule encoding a self assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle. The retroviral particle comprises a modified HIV genome devoid of long terminal repeats containing a nucleotide sequence coding for a chimeric envelope glycoprotein which has a first (a) and second (b) retroviral envelope amino acid sequence, where (a) contains the HIV-1 gpl20 conserved region 2, and (b) contains a retroviral envelope amino acid sequence of a heterologous strain of HIV-1, HIV-2, HIV-I or HILV-II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Retrovirus-like particle, modified HIV genome;
chimeric envelope glycoprotein; HIV-1 gp120; conserved region 2; HIV-1;
HIV-2; HTLV-1; HTLV-II; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunogenic retrovirus like particles comprising modified HIV genomes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 77; DB 2; Length 21; 100.0%; Pred. No. 0.00012; ive 0; Mismatches 0; Indels
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Example 4; Col 7-8; 12pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY16052 standard; peptide; 21 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       useful as vaccines against HIV.
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(first entry)
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                            Sequence 21 AA;
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20-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a Human immunodeficiency virus type 1 derived peptide. The peptide is used in the immunoassay diagnostic kit of the invention. The specification describes an immunoassay diagnostic kit for detecting antibodies in a sample, which comprises an antigen consisting of a self-assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle encoded by a modified HIV genome that is devoid of long terminal repeats and contains a nucleotide sequence coding for a chimeric envelope protein having a first amino acid sequence containing HIV-1 gpl20 conserved region 2 and a second amino acid sequence containing an envelope sequence of a heterologous strain of HIV-1, HIV-2, HIUV-1 or HTUV-11. (Updated on 20-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunoassay diagnostic kit for detecting antibodies - comprising chimeric retrovirus-like particles.
inserted into (a) at an endogenous site (BglII and StuI). (b) may comprise peptides AAY16049-51 and AAY16055. The nucleic acids are useful as vaccines against HIV. The present sequence is used in the course of the invention. (Updated on 20-MAR-2003 to correct PR field.) (Updated on 17-0CT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunoassay diagnostic kit, antibody detection;
chimeric envelope protein; HIV-1 gpl20 conserved region 2; HIV-1; HIV-2;
HTLV-1; HTLV-11.
                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ch 100.0%; Score 77; DB 2; Length 21; 1 Similarity 100.0%; Pred. No. 0.00012; 15; Conservative 0; Mismatcher
                                                                                                                               Length 21;
                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1 derived peptide.
                                                                                                                             Score 77; DB 2; I
Pred. No. 0.00012;
                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rovinski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 4; Col 9-10; 12pp; English.
                                                                                                                                                                                                                                                                                                          AAW85568 standard; peptide; 21 AA
                                                                                                                                                             o;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus 1
                                                                                                                             100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95US-00452503
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93US-00073526
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(first entry)
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                                                                                                                            Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                           7 RIQRGPGRAFVTIGK
                                                                                                                                                                                            1 RIQRGPGRAFVTIGK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-069713/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Klein MH, Cao SX,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 AA;
                                                                                                Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                       20-MAR-2003
24-FEB-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                         AAW85568;
                                                                                                                                                                                                                                                                            RESULT 65
   8X38888
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The present invention relates to diphosphonate compounds useful as immunological adjuvants. The compounds can be used for stimulating an immune response to an antigen. The present sequence is an immunogeneic peptide used to test the ability of the compounds to cause an increase in the immune response. The peptide consists of an amino terminal cysteine residue, a glycine/alanine/glycine spacer and amino acids 308-322 of the V3 loop of HIV IIIB gp120 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Retrovirus-like particle CLTB73 containing a V3 (HXB2) antigenic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus; HIV; retroviral antigen; gag; pol; env; immune response; antigenic marker; antigenic epitope; retrovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                        ن
                                                                                                                                                                                                                                                                                                                                                                                                                                      diphosphonate compounds, useful as immunological adjuvants for
                                                                                                                                                                                                                                                                                                                                                                        Rose
                                                                                                                                                                 Peptide P18 derived from V3 loop of HIV IIIB group 120 protein.
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                                                                                                                                                                                                                                                                                                                                                                        Nault A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 21;
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                                                                                                                                                                                                                                                                                                                                                                        Mcguiness P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 77; DB 3; I 100.0%; Pred. No. 0.00012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    stimulating an immune response to an antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                        Lewis M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 8; Page 86; 130pp; English.
                                                                                     AAB15012 standard; peptide; 21 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus.
                                                                                                                                                                                                                                                                                               01-FEB-2000; 2000WO-US002755
                                                                                                                                                                                                                    Human immunodeficiency virus
12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIORGPGRAFVTIGK 15
             RIORGPGRAFVTIGK 20
                                                                                                                                                                                           HIV; immune; diphosphonate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                        Ishizaka ST,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100.
RIORGPGRAFVTIGK
                                                                                                                                                                                                                                                                                                                                               (EISA ) EISAI CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-514809/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 AA;
                                                                                                                                                                                                                                            WO200044758-A1
                                                                                                                                                                                                                                                                                                                                                                                       Rossignol DP;
                                                                                                                                                                                                                                                                                                                        01-FEB-1999;
                                                                                                                                        07-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                          Hawkins LD,
                                                                                                                AAB15012;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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AAU08699
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                                                                            AAB15012
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Gaps

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Query Match Best Local Similarity Matches 15; Conserv

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WPI; 1994-263787/32.
                                                                                                                                                                                                                                       Sequence 22 AA;
               01-APR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JAN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
21-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-AUG-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUL-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-1993;
21-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                          Emini EA,
                                                                                                                                                                                                                                                                                                                                                                                                          AAR57470;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AIDS; 88.
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(WANG/)
(CONE/)
(MERV/)
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                                                                                                                                                                                                                                                                                                                                                              RESULT 69
                                                                                                                                                                                                                                                                                                                                                                         AAR57470
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                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a method for determining the presence of sample antibodies specifically reactive with HIV retroviral antigens in a sample. This involves contacting a sample suspected of containing HIV-specific antibodies with a non-infectious, non-replicating, immunogenic HIV-like particle as an antigen. The antigen comprises an assembly of a gag gene product, a pol gene product and a modified env gene product containing a non-retroviral heterologous, antigenic, anchor sequence that replaces the endogenous anchoring functions of the env gene product. The method detects immune complex formation between HIV-specific antibodies and the antigens. The method is also useful for identifying antiserum enthod entered by immunisation with an immunogenic composition capable of eliciting HIV-specific immune response. The antigenic marker may comprise at least one antigenic epitope from another virus. This sequence represents a retrovirus-like particle containing an antigenic marker
                                                                                                                                                                           Differentiating between infection by human immunodeficiency virus (HIV) and antiserum generated by immunization against HIV, comprises use of non-infectious, non-replicating HIV-like particle with heterologous, antigenic anchor sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Immunodeficiency Virus; antigen; ELISA; recombinant antibody; throneutralising monoclonal antibody; immunoglobulin; AIDS; acquired immune deficiency syndrome; chimeric antibody; surface glycoprotein gp120; V3 loop; epitope mapping.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 4; Length 21; 100.0%; Pred. No. 0.00012; ive 0; Mismatches 0; Indels
                                                                                                                                    Persson R, Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1; (IIIB isolate).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gp120 V3 loop sequence of HIV-1 IIIB isolate.
                                                                                                                                                                                                                                  Disclosure; Col 17; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR42153 standard; peptide; 22 AA.
                                                                 98US-00027955
                                                                                       98US-00027955
                                                                                                                                    Yao F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93WO-US002629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(revised)
(first entry)
                                                                                                            (CONN-) CONNAUGHT LAB LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15; Conservative
                                                                                                                                  Rovinski B, Cao S,
                                                                                                                                                      WPI; 2001-595518/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 AA;
                      US6291157-B1
                                                                                       23-FEB-1998;
                                                                 23-FEB-1998;
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25-MAR-2003
27-APR-1994
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                                           18-SEP-2001
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 Synthetic.
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Matches
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Antibodies able to neutralise more than one HIV-1 isolate are claimed. The gpl20 V3 loop sequences from different isolates comprising the Principal Neutralising Determinant motif GPGR are given in AAR42153-R42161. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                                                               New recombinant human antibody - with HIV neutralising activity against at least two isolates, useful for preventing or treating infection in diagnosis, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunisation; vaccine; therapy; prophylaxis; defective gene;
non-functional gene; template; antisense; ribozyme; bupivacaine;
human immunodeficiency virus; acquired immune deficiency syndrome; HIV;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 22;
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                                                                                          Pfarr DS;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 77; DB 2; Lu
100.0%; Pred. No. 0.00013;
Mismatches 0;
                                                                                            Johnson LS,
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                                                                                                                                                                                                                                                                         Example 16; Page 100; 154pp; English.
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                                                                                          Mark GE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR57470 standard; protein; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93US-00008342.
93US-00029336.
93US-00093235.
93US-00124962.
93US-00125012.
92US-00861701
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 RIORGPGRAFVTIGK 15
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV BRU V3 loop peptide.
                                          (MERI ) MERCK & CO INC.
                                                                                          Conley AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WEINER D B.
WILLIAMS W V.
WANG B.
                                                                                                                                    WPI; 1993-336600/42.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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treatment of AIDS. (Updated on 16-OCT-2003 to standardise OS field)
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ID AAY8
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                                                                                    the sequence encoding against HIV contains a DNA construct which comprises the sequence encoding apple0. The genetic material was then introduced into the cells of an individual by (a) contacting the individual's cells with a polymucleotide function enhancer (bupivacaine) and (b) administering to the cells the nucleic acid molecule free of retroviral particles. Wucleic acid molecules which are delivered to cells may serve as genetic templates for proteins that function as prophylactic and/or therapeutic immunising agents; replacement copies of defective, missing or non-functional genes; genetic templates for therapeutic proteins; or non-functional genes; genetic templates for the sequence of contact templates for antisense molecules or as genetic templates for continuous and defective, missing or non-functional genes; genetic templates for antisense molecules or as genetic templates for contact by HIV neutralising antibodies) and was used to epitope targetted by HIV neutralising antibodies and was used to determine whether the anti-gpi60 antibodies elicited in mice immunised with the genetic vaccine were reactive with this region. (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This is the sequence of the V3 loop of the gp120 envelop protein from HIV -1 strain IIIb. The sequence was used with a construct comprising part of the HIV-1 gp120 V3 loop common consensus PND sequence (AAMV07391) fused to part of the HIV-1 CD4 binding shie (AAMV07391) and with the V3 loop sequences from HIV-1 strains Thai B (AAMV07393) or HGP-30 (AAMV07394) to generate a lysine branched peptide which is useful for the prevention and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               wovel AIDS vaccine - comprises branched lysine peptide fragments derived from HIV env protein.
Method for introducing genetic material into cells - utilises polynucleotide function enhancer and nucleic acid free of retroviral particles, e.g. HIV immunisation.
                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV-1; gpl20; V3 loop; common consensus PND domain; envelop; CD4; binding site; stem-loop; lysine branched peptide; AIDS.
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0
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                                                             Example 3; Page 44; 136pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 5-6; 8pp; Japanese.
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Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       8 RIORGPGRAFVTIGK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (OKUD/) OKUDA K.
                                                                                                                                                                                                                                                                                                                                    Sequence 22 AA;
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24-FEB-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence represents the V3 loop from the gpl20 protein of the human immunodeficiency virus-1 (HIV-1) strain IIIB. The invention relates to the generation of heterohybridomas producing human monoclonal antibodies (see AAX79204-X79207) to a neutralising epitope of HIV-1 prepared by transforming peripheral blood lymphocytes with Epstein-Barr virus. The antibodies can be used to treat someone infected with HIV-1 or suffering from AIDS. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Light chain; variable region; human; HIV-1; gp120; monoclonal antibody; epitope; V3 loop; heterohybridoma; human immunodeficiency virus-1; peripheral blood lymphocyte; Epstein-Barr virus; EBV; AIDS.
                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Heterohybridoma producing human monoclonal antibodies to human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 77; DB 2; Length 22; 100.0%; Pred. No. 0.00013; ive 0; Mismatches 0; Indels
                                           Length 22
                                                                                          0; Indels
                                           Score 77; DB 2; 1 Pred. No. 0.00013;
                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV-1 strain IIIB gp120 V3 loop sequence.
                                                                                                                                                                                                                                                                                                    AAY07488 standard; peptide; 22 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY85137 standard; protein; 22 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus 1.
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91US-00684090.
92US-00872675.
                                              100.0%;
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                 Query Match
Best Local Similarity 10v...
Best 15; Conservative
                                                                                                                                        1 RIORGPGRAFVTIGK 15
                                                                                                                                                                               RIORGPGRAFVTIGK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gorny MK, Zolla-Pazner S;
                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  [mmunodeficiency virus-1
                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-370481/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 AA;
Sequence 22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-JUN-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                 17-OCT-2003
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Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        loop peptide sequence. The prepride sequence is used to test the reactivity of the antibodies of the invention. The invention relates to a method for the production of neutralising antibodies against conformational epitopes of HIV-1 envelope proteins in humans. The method comprises administering to a human, a recombinant uncleaved gp140 protein retaining its oligomeric structure. The human produces neutralising antibodies against conformational epitopes of the HIV-1 gp140 protein antibodies of the invention can be used for preventing and diagnosing an antibodies of the invention can be used for preventing and diagnosing an HIV infection. A diagnostic method using the antibodies involves is labelled monoclonal antibody for gp140, and centacting it with a labelled monoclonal antibody for gp140, and detecting any bound antibody. (Updated on 12-SEP-2003 to standardise OS field)
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                                                                                    Human immunodeficiency virus type 1; HIV-1; infection; prevent; detect; glycoprotein 140; gp140; neutralising antibody; conformational epitope; V3 loop.
                                                                                                                                                                                                                                                                                                                                                                          This sequence represents a human immunodeficiency virus type-1 IIIB V3-
                                                                                                                                                                                                                                                                                                        Producing neutralizing antibodies useful for preventing, treating and diagnosing an HIV infection in a mammal comprises administering recombinant uncleaved gpl40 proteins to a human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human N-acetylgalactosaminyl transferase T4, GalNac T4, substrate #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GalNac T4; N-acetylgalactosaminyl transferase T4; acceptor substrate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 77; DB 3; Length 22; 100.0%; Pred. No. 0.00013;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                          (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                 Broder CC;
                                                              HIV-1 IIIB V3 loop peptide sequence.
                                                                                                                                                                                                                                                                                                                                                    Example 10; Col 12; 15pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABU07537 standard; peptide; 22 AA.
                                                                                                                                 Human immunodeficiency virus 1.
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                                                                                                                                                                                                97US-00805889
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                                                                                                                                                                                                                                                                 Earl PL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIORGPGRAFVTIGK 15
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                                           (first entry)
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 RIORGPGRAFVTICK
                                 (revised)
                                                                                                                                                                                                                                                                                     WPI; 2000-270121/23.
                                                                                                                                                                                                                                                                Doms RW, Moss B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 AA;
                                                                                                                                                                                              03-MAR-1997;
                                                                                                                                                                                                                    10-DEC-1993;
                                12-SEP-2003
20-JUN-2000
                                                                                                                                                    US6039957-A
                                                                                                                                                                           21-MAR-2000
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13-MAR-2003
           AAY85137;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 73
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The invention relates to glycosylating a MUC1 (mucin 1) acceptor substrate, comprising glycosylating the substrate with enzymatically active N-acceptyglaterosaminyltransferase (GalNRc)-T1, GalNAc-T2 or GalNAc-T3, or with GalNAc capable of glycosylating MUC1 glycosylation sites that can be glycosylated by GalNAc-T1, GalNAc-T2 or GalNAc-T3, and glycosylated pubstrates with enzymatically active human GalNAc-T4 to glycosylate specific Ser, Thr positions in the MUC1 substrate. The method is used for glycosylating an MUC1 acceptor substrate. The method substrates are useful in preparation of vaccines and antinflammatory substrates are useful in preparation of vaccines and antinflammatory previously characterised GalNAc transferases. The activity of GalNAc-T4 exhibite a different substrate specificity than preparation of vaccines and antinflammatory is unique and specific to glycosylate specific serine and threonine is unique and specific to glycosylate specific serine and threonine substrate peptide used to test the substrate specificity the human GalNac T4 protein, HIVHIBGPI20. (Updated on 23-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Glycosylating MUC1 acceptor substrate, by glycosylating substrate with N-acetylgalactosaminyltransferase T1, GalNAc-T2 or GalNAc-T3, then with human GalNAc-T4 to glycosylate specific Ser, Thr residues in substrate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV; therapy; AIDS; principal neutralising domain; antibodies; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cpd. eliciting, binding with neutralising antibodies to HIV variants.
glycosylation; mucin 1; MUC1; vaccine; antiinflammatory; GalNAc-T1;
GalNAc-T2; GalNAc-T3; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaetgens J;
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100.0%; Score 77; DB 6; Length 22;
Best Local Similarity 100.0%; Pred. No. 0.00013;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eisenkraetzer D,
                                                                                                                                                                                                                                                        /label= OTHER
/note= "Gln is acetylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bennett EP,
                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; Col 9; 10pp; English
                                                                                                           Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hassan FH, Clausen H,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GLYC-) GLYCOZYM APS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-147066/14.
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                                                                                                                                                                                   Key
Modified-Bite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-DEC-1998;
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25-MAR-2003
:20-SEP-1990
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The present invention relates to an immunogenic composition comprising a recombinant uncleaved gp140 protein which is a C-terminally truncated form of HIV-1 gp160 protein, missing the gp41 transmembrane domain, and retaining its oligomeric structure, such that neutralizing antibodies against conformational spitopes of HIV-1 envelope proteins found on the oligomeric structure of are produced in an immunized human. The invention is useful for stimulating an anti-HIV-1 env immune response in a mammal, by stimulating the formation of neutralizing antibodies against conformational epitopes of HIV-1 env protein in a mammal. gp140 is also useful for preventing HIV infection in a mammal. (Updated on 11-SBP-2003 to standardise OS field)
                                      Immunogenic composition for stimulating mammalian immune response, comprises recombinant uncleaved gp140 protein retaining its oligomeric structure such that antibodies against HIV envelope proteins are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protease inhibitor may be use to suppress organ transplant rejection without serious side effects. It may also be used in prevention and therapy of allergy, aplastic anaemia and systemic erythaematodes. See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protease inhibiting peptide immuno-suppressant - used to suppress rejection reaction in organs transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                autoimmune disease; allergy; aplastic anaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2; Length 24;
                                                                                                                                                                                                                                                                                                                                         Length 23;
                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                         100.0%; Score 77; DB 4; I
100.0%; Pred. No. 0.00013;
iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunosuppressant protease inhibitor.
                                                                                                             Example 10; Col 13; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR06211 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 181; 6pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88JP-00310635.
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                                                                                                                                                                                                                                                                                                                                         Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               systemic erythaematodes.
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           WPI; 2001-167730/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Organ transplant;
                                                                                                                                                                                                                                                                                                                Sequence 23 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                  produced
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                                                                                                                                                                                                                                                                                                                                         Either the N-terminal (a) or C-terminal (b), but not both, may be omitted 'ether (a) or (b) may comprise any of the following: cysteine, a protein or other moiety capable of enhancing immunogenecity, a peptide from an HIV principal neutralising domain, peptide capable of stimulating T-cells, or general immune stimulant. See also AAR04427-R04506 and AAR04437-R04506 and on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR Field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Revised record issued on 09-SEP-2004 : Correction to Peature Table Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                   Principal neutralising domain of HIV variants - used for producing peptide (8) and antibodies for diagnosis, prophylaxis and/or therapy therapy of HIV infection.
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                                                                                                                                                                                  Grimaila R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 77; DB 2; Length 23; 100.0%; Pred. No. 0.00013;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                  Farley J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV; Human immunodeficiency virus; immune; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Moss B;
                                                                                                                                                                                  Javaherian K,
                                                                                                                                                                                                                                                                                                               Claim 27 (d); Page 84; 108pp; English.
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                                                                                             88US-00252949.
89US-00359543.
89US-00407663.
                                                                    88US-00252949.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15; Conservative
                                                                                                                                                                                  Rusche JR, Putney SD,
Lynn DU, Petrobre J;
                                                                                                                                                      (REPK ) REPLIGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Broder CC,
                                                                                                                                                                                                                            WPI; 1990-147824/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-DEC-1993;
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                                                                    03-OCT-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-APR-1998;
                                                                                                03-OCT-1988;
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09-APR-2001
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              WO9003984-A.
                                                                                                             01-JUN-1989
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                                         19-APR-1990
                                                                                                                                                                                 Rusche JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                            PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB66704;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 75
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Gaps

Di Bello C;

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The principal neutralizing domain (PND) of HIV-1 corresp. to a 24- amino acid sequence arranged in a loop determined by a disulfide bridge in the third hypervariable region, '03, of the protein gp 120. The central portion of the V3-PND contains a sequence which is highly conserved in different HIV-1 isolated strains, whereas the amino acids flanking this sequence are variable. The antigenic properties of V3 region are known to be virus-specific, antibodies elicited by MN-derived peptide do not neutralize HTLV-III B virus and vice-versa. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      V3 loop; gp41; envelope protein; MN; prototype; virus; varient; HIV; homology; heteroconjugate; enzyme; epitope mapping; replication; conjugate; immunogenic carrier; keyhole limpet hemocyanin; KLH; ovalbumin; succinyl maleimidomethyl cyclohexanylcarboxylate; SMCC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic peptide(s) which enhance infectivity of HIV-1 in cellular cultures - are used for determining HIV-1 virus in blood and other biological materials.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Not in the natural sequence of this isolate"
                    Diagnostic; assay; detection; AIDS; human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 0.00014;
                                                                                                                                                                                                                                                                                                                                                      Mammano F, Panozzo M, Dettin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                               Human immunodeficiency virus 1; strain HTLV-III B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Heteroconjugate antibody immunogen RP135 (IIIB).
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                                                                                                                                                                                                                                                                                 (SUPE-) INST SUPERIORE DI SANITA.
(CNDR ) CONSIGLIO NAZ DELLE RICERCHE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR29233 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 17; 31pp; English.
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(first entry)
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Best Local Similarity 100.0
                                                                                                                                                                                                                                                                                                                                                 Pasti M,
                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-299983/36.
                                                                                                                                                                                                                                                                                                                                                 De Rossi A, Past:
Chieco-Bianchi L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 24 AA;
                                                                                                                                                                                               29-JAN-1992;
                                                                                                                                                                                                                                           30-JAN-1991;
                                                                                                           WO9213882-A1
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14-APR-1993
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                                                                                                                                                      20-AUG-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR29233;
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                      셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           XHXLLHXSXEEEXBXBXHLLHXSX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prepn. of anti-HIV sulphated polysaccharide - by affinity chromatography using a resin-bound peptide corresp. to a HIV gp. 120 fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-HIV sulphated polysaccharide (SPS) can prevent syncytium formation in HIV infected C4 cells. SPS may be isolated by affinity chromatography with the given resin bound peptide fragment RP135. (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                      Residues 301-324 of HIV gp 120 protein used in isolation of sulphated polysaccharide by affinity chromatography.
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                                         Gaps
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                                         0; Indels
               Pred. No. 0.00014;
                                         Mismatches
                                                                                                                                                                                                                                    AAR07018 standard; peptide; 24 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR26565 standard; peptide; 24 AA.
100.08; Pr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RICH ) MERRELL DOW PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV; AIDS; ARC; gp120; RP135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90CA-02007258.
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                                                                                                           8 RIQRGPGRAFVTIGK 22
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(revised)
(first entry)
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(first entry)
               Best Local Similarity 100.
Matches 15; Conservative
                                                                               1 RIQRGPGRAFVTIGK
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1990-290631/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 AA;
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18-JAN-1991
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28-JAN-1993
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ID AARO7018

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DY 28-C
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The sequence is that of peptide IIIB, derived from the V3 region of HIV gp120, it was used in binding assays for anti-gp120 antibodies. The anti-gp120 antibodies are useful in vaccine formulations for the treatment or prevention of HIV infection. See also AAR26867-R26973. (Updated on 25-MAR-2003 to correct PN field.)
          Sxample, Page 9; 30pp; Japanese.
                                                                                                                                   1 Similarity 100.
                                                                                                  Sequence 24 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 AA;
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04-JUL-1993
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                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                        Query Match
Best Local S
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                                                                                                                                              Matches
                                                                                                                                                                                                                            RESULT 81
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                                                                                                                                                                                               The sequences given in AAR29226-15 represent peptides which were used as immunogens for the production of antibodies against HIV. These peptides may be either unconjugated or conjugated to an immunogenic carrier, eg. a keyhole limpet hemocyanin (KIH) or ovalbumin, using succinyl maleimidomethyl cyclohexanylcarboxylate (SMCC) as a conjugating agent. Viruses containing these or similar sequences may be recognised by the heteroconjugate enzymes of the invention. The antibodies raised against these sequences may be identified by standard epitope mapping techniques. These autibodies are capable, even at low concentrations, of nearly eliminating viral replication of different strains of HIV. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                Hetero-conjugate antibodies for treating HIV infections - comprise an antibody specific for an effector cell surface antigen and an antibody to a V3 loop of GP-120 envelope protein of HIV.
                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-idiotype antibodies and methods for their selection - useful as vaccines for the prevention and treatment of HIV infection.
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                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 0.00014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus; AIDS; anti-gp120 antibodies.
                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV gp120 V3 region binding assay peptide IIIB.
                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                              Disclosure, Page 19; 69pp; English
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                       92WO-US003616.
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                                            91US-00699773
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(first entry)
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Matches 15; Conservative
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                                                                                        Higgins PJ, Potts BJ;
                                                                  (REPK ) REPLIGEN CORP
                                                                                                             WPI; 1992-415475/50.
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                                                                                                                                                                                                                                                                                                                                             Sequence 24 AA;
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06-MAR-1992;
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                                            14-MAY-1991;
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                       29-APR-1992;
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20-MAY-1998
26-NOV-1992
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Nine peptides from the V3 regions of HIV-1 isolates IIIb, RF and MN were incorporated into tetravalent multiple antigen peptide systems (MAPS) (Ree AAR33406-44). Parallel groups of three peptides with chain lengths spanning from 11-24 residues were synthesised in MAPS formate for each isolate. BLIS assays demonstrated that antisera titers in mice were immunisation to the length of the IIIb peptide used for the immunisation - the longer the stronger the response. There was no substantial antibody prodn. in mice against the other two series of peptides, RF (B4-B6), and MN (B7-B9), except for a low reactivity in the gpo immunised with B8 (MN isolate). Specificity tests of the B cell response demonstrated that the T cell epitope (AAR32415) also serves as a B cell epitope. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence of peptide B1 which comprises AAs 308-331 from the V3 region of HIV-1 isolate IIIb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New multiple antigen peptide(s) as HIV vaccines - include a dendritic core covalently bonded to peptide including the sequence IGFGR.
100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 0.00014; Pred. No. 0.10018; ive 0; Mismatches 0; Indels
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Pred. No. 0.00014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1; vaccine; dendritic core; ss
                                                                                                                                                                                                                                                                                                                                                                                                                 AAR32406 standard; peptide; 24 AA.
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                                                                                                                                                       1 RIORGPGRAFVTIGK 15
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(first entry)
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(UYRQ ) UNIV ROCKEFELLER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1993-093730/11.
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Matches

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HIV gpl20 V3 loop-derived peptides (AAR38170-87) are successful in generating CTL responses, esp. peptide R15K (AAR38187); the T-helper cell-inducing peptide includes the sequence C19A (AAR38187); the T-helper cell-inhibiting peptides are given in AAR38188-206, and are esp. peptides R15K, N24G, E13V, R8K, T13Q and H13N (AAR38165-69). The peptides may also be derived from an influenza virus protein or a sendai virus protein (AAR41014-15). It was observed that peptide N24G (amino acids 308-311), with sequences derived from the V3 loop of H1V-1 IIIB, inhibits H1V-1 infection of primary human T cells by 92% at 1 microg/m1 (ca. 0.4-0.6 microW). (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG
                                                                                                                                                                                                                                                                                                                                                                    Peptide composition for treating and preventing viral infections - comprise CTL-inducing epitope and HIV infection-inhibiting sequence or helper cell-inducing sequence.
                      gp120; HIV-1; cytotoxic T-lymphocyte; CTL; T-helper; AIDS; infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antigen; B2; third variable domain; V3 loop; gp120; HIV-1; vaccine; strain IIIB; multiple antigenic peptide system; dendritic core; lipophilic membrane anchoring group; mammal; humoral; immunisation; cytotoxic T cell; CT; immune response; infection; Freunds adjuvant; pathogen; HIV; influenza; malaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gp120 V3 loop antigen B2 and lipophilic membrane anchoring group.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 0.00014; Pred. o. Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                   Nehete PN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1. .18
/label= B2 antigenic peptide
19. .24
                                                                                                                                                                                                                                                                                                 Platsoucas CD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 19; Page 95; 130pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR44191 standard; peptide; 24 AA.
                                                         Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1
                                                                                                                                                                                                        91US-00800932.
92US-00945865.
                                                                                                                                                                                                                                                                                                 Sastry JK, Arlinghaus RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RIORGPGRAFVTIGK 22
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(first entry)
                                                                                                                                                                                                                                                             (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15; Conservative
                                                                                                                                                                                                                                                                                                                                    WPI; 1993-196739/24.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 AA;
                                                                                             WO9310816-A1
                                                                                                                                                                     02-DEC-1992;
                                                                                                                                                                                                        02-DEC-1991;
                                                                                                                                                                                                                         16-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
20-MAY-1994
                                                                                                                                 10-JUN-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR44191;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Two peptides, gp41[8] and V3 loop derived from the gp120 region of HIV-1 were synthetised and purified. To improve the solubility of the gp41[8] peptide the sequence RSS was added to the amino terminal to produce peptide R-S-Sgp41[8]. The immunodominant HIV-derived peptides were used to ascertain whether E.coli outer membrance protein TraT augments the in vitro T-cell proliferative responses. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Compans. contg. E.coli outer membrane proteins TraT, OmpA or OmpF -
increase immune response and are used for treating auto:immune diseases,
   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 0.00014;
   0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bell SJD, Cooper DA
                                                                                                                                                                                                                                                                                                                              AIDS; HIV; therapy; autoimmune disease; gp120; ss.
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                          Sequence of HIV-1 derived V3 loop peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BIOT-) BIOTECH AUSTRALIA PTY LTD.
(SVIN-) ST VINCENT'S HOSPITAL SYDNEY LTD.
                                                                                                                                                              AAR33190 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR38165 standard; peptide; 24 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 91AU-00007725
                                                         1 RIORGPGRAFVTIGK 15
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(revised)
(first entry)
                                                                                                                                                                                                                                        (revised)
(first entry)
15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1993-093727/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AIDS, cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 24 AA;
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25-MAR-2003
12-OCT-1993
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                                                                                                                                                                                                                                                                                                                                                                                                      WO9303762-A1
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11-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                  AAR33190;
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Matches

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RESULT 83
AAR38165
AD AAR38165
XX
XX
AC AAR38
DT 27-AU
DT 25-MA
DT 12-OC
XX
XX
XX
DT 25-MA

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Gaps

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WPI; 1998-556461/47.
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                                                WPI; 1994-341488/42
                                                                                                                                                                                                                              Local Similarity
nes 15; Conserv
                                                                                                                                                                                                    Sequence 24 AA;
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                              Vahlne A,
                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                       AAW67414;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chong P,
                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                             V3 loop
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                                                                                                                                                                                                                                                                                                                RESULT 86
                                                                                                                                                                                                                                                                                                                          AAW67414
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                                                                                                                                                                                                                    The sequence given in AAR44190 is a peptide antigen, B2, which represents residues 312-329 of the third variable domain (V3 loop) of gp120, of HIV-1 strain IIIB. This sequence was attended to an aminon acid linker (see also AAR4191) in the production of a multiple antigenic peptide system. This system comprises a dendritic core to which are covalently attached at least one peptide, eg. an antigenic peptide, and a lipophilic membrane concloring group. This system may be injected into a mammal and elicits both humoral and cytotoxic T cell (CTL) immune responses. This system may be unchoring group provides efficient adjuvant activity without the toxicity problems of Freunds adjuvant, while the dendritic structure allows multiple antigens to be attached. Optionally the antigens may be derived than one disease, eg. HIV, influenza and malaria. (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1; HIV-1; gpl20 epitopes; vaccines;
HIV neutralising antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                          New multiple antigen system esp, for use in HIV vaccines - contains lipophilic membrane anchor imparting adjuvant activity, and peptide antigens coupled to dendritic core.
                                                                                                                                                                                                                                                                                                                                                                                                                            ..
 /note= "Lipophilic membrane anchoring group"
                                                                                                                                                                                                                                                                                                                                                                                                       Score 77; DB 2; Length 24; Pred. No. 0.00014; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV-1 gp120-24 epitope amino acids 307-330.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR63821 standard; peptide; 24 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                              92US-00877613
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                                                           93WO-US004179
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                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 RIORGPGRAFVTICK 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                 (UYRQ ) UNIV ROCKEFELLER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
                                                                                                                                          WPI; 1993-368723/46.
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24 AA;
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25-MAR-2003
29-JUN-1995
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                                                           03-MAY-1993;
                                                                              01-MAY-1992;
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                    WO9322343-A1
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                                        11-NOV-1993
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                                                                                                                       ram JP;
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AAK63809-R63849 are epitopes from the human immunodificiency virus type 1 (HIV-1) gp120, by binding one or more of these epitopes to a carrier a HIV vaccine is produced. These vaccines can elicit the production of HIV-neutralising antibodies in monkeys, and therefore may be used to prevent HIV infections, and to highten the immune response in HIV infected humans. (Updated on 25-MRR-2003 to correct PN field.) (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunogen; vaccine; HIV-1; T-cell; B-cell; epitope; core protein; gp120;
                                                                                                                                                                                      New peptide(s) comprising HIV gp120 epitope(s) - for prodn. of vaccines against HIV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic human immunodeficiency virus-1 peptide(s) - containing T-cellepitope and B-cell epitope(s) are candidate vaccines against HIV-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-cell epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitopes are generally designed based on the p24 core protein and the B-cell epitopes from the V3 loop of the gpl20 protein from various HIV-1 strains. This sequence corresponds to an HIV-1 B-cell peptide epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                 Horal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 77; DB 2; I 100.0%; Pred. No. 0.00014;
                                                                 Jeansson S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                 Rymo L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW67414 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 3; 40pp; English
                                                                                                                                                                                                                                                                                                 Claim 1; Page 18; 77pp; English.
(SYNT-) SYNTELLO VACCINE DEV AB
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94US-00257528.
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                                                                 Svennerholm B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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(revised)
(first entry)

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Enhancing the immune response using a recombinant human low-density lipoprotein receptor, useful for treating viral infections, especially human immunodeficiency virus (HIV) infection.
                                                                                                                         immunodeficiency; neoplastic tissue; myalgic encephalomyelitis; ME; viral infection fatigue syndrome; tuberculosis; hepatitis; AIDS; ARC; acquired immunodeficiency syndrome; AIDS related complex; HIV-infected CD4 cell; immunosuppressive peptide.
                                                                                                              HIV; LDL; low density lipoprotein; human; immune response; infection;
                                                                                       HIV LDL binding peptide, sequence A.
   AAY22581 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 7; Page 19; 24pp; English.
                                                                                                                                                                                       Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                           (PREN/) PRENDERGAST P T.
                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-494040/41.
                                                                                                                                                                                                                                                                                                                                    Prendergast PT;
                                                                                                                                                                                                              WO9938524-A2.
                                                                                                                                                                                                                                                             28-JAN-1999;
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                                                   17-OCT-2003
19-OCT-1999
                                                                                                                                                                                                                                      05-AUG-1999.
                            AAY22581;
   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a synthetic peptide comprising an amino acid sequence containing a T-cell epitope of an HIV gag protein linked a its C terminus to an amino acid sequence containing a B-cell epitope of an HIV gp41 protein and containing the amino acid sequence: XILKDWX2; where XI = E, A, G or Q, and X2 = A or T, or an amino acid sequence capable of aliciting an HIV-epecific antiserum and recognizing the sequence XILKDWX2. The synthetic peptide is useful in vaccines against HIV infection and in diagnostic applications. AAW98892 to AAW98896, and AAW998899 to AAW998996, and
                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic chimeric HIV polypeptides - comprising gag protein T-cell epitope linked to gp41 B-cell epitope.
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                                                                                                                                                                                                                                                                               HIV-1; human immunodeficiency virus, vaccine, T-cell epitope, gag protein; B-cell epitope, gp41 protein; chimeric; infection.
                                                 100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 0.00014;
                                                                       0; Indels
                                                                                                                                                                                                                                                        HIV-1 vaccine synthetic peptide SEQ ID NO:99.
                                                                          0; Mismatches
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                                                                                                                                                                                AAW98904 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1.
used to immunise a guinea pig
                                                                                                                                                                                                                                                                                                                                                                                                                               93US-00073378.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chong P, Klein MH, Sia CDY;
                                                                                                                                                                                                                                                                                                                                                                                                        95US-00462507
                                                                                                1 RIORGPGRAFVTIGK 15
                                                                                                                     7 RIORGPGRAFVTIGK 21
                                   Query Match
Best Local Similarity 100...
Thea 15, Conservative
                                                                                                                                                                                                                                05-MAY-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CONN-) CONNAUGHT LAB LTD.
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                        Sequence 24 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                      05-JUN-1995;
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09-JUN-1994;
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                                                                                                                                                                                                                                                                                                                                                                              02-MAR-1999
                                                                                                                                                                                                                                                                                                                    Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention
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98US-0072980P. 99WO-IB000149.

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This sequence represents a HIV sequence that binds human low density lipoprotein (LDL), and is designated sequence "A". The invention relates to a method for enhancing the immune response in a patient with a condition, selected from immunodeficiency (due to a viral, bacterial, mycoplasmic, fungal or parasitic infection, or from the growth of mycoplasmic, fungal or parasitic infection, or from the growth of comprises using a pharmaceutical composition, comprising a recombinant comprises using a pharmaceutical composition, comprising a recombinant comprises using a pharmaceutical composition, comprising a recombinant compositions for: the treatment of acquired immunodeficiency syndrome compositions for: the treatment of acquired immunodeficiency syndrome compositions for: the treatment of acquired immunodeficiency syndrome compositions for: the treating blood or body fluid or organs to neutralise/remove immunosuppressive peptides and/or viruses; or treating infection in a human or animal host. The human recombinant LDL receptor is also useful for manufacturing medicaments for treating all the conditions given above. The human recombinant LDL receptor conditions given above. The human recombinant LDL receptor is conditions given above. The human recombinant LDL receptor is conditions given above. The human recombinant LDL receptor is conditions given above. The human recombinant LDL receptor is conditions given above. The human recombinant LDL receptor is to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15; Conservative
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Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 24 AA;
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ID AAY2
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Gaps

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0; Indels

Mismatches

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15; Conservative

Matches

Query Match Best Local Similarity

RESULT 88 AAY22581

Length 24;

100.0%; Score 77; DB 2; I 100.0%; Pred. No. 0.00014;

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This sequence represents a variant of the HIV sequence that binds human low density lipoprotein (LDL), and is designated sequence "A" (see AAV22581). The sequence "A" peptide is solated from HIV isolate IIBB(BHO), and this sequence was isolated from HIV isolate IIBB(BHO). The invention relates to a method for enhancing the immune response in a patient with a condition, selected from immunodeficiacy (due to a viral, bacterial mycoplasmic, fungal or parasitic infection, or from the growth of neoplastic tissue), myalgic encephalomyelitis (ME), post inoculation or viral infection fatigue syndrome, tuberculosis, or hepatitis. The method comprises using a pharmaceutical composition, comprising a recombinant human LDL receptor or a mimic molecule to the cystein e rich domain of LDL receptor. The human recombinant LDL receptor forms commissions where the complexitions of the compositions of the composition of LDL receptor forms financeutical compositions for RC (ALDS related complex); reducing syndrome (ALDS) or ARC (ALDS related complex); reducing syndrome (ALDS) or ARC (ALDS related complex); reducing syndrome (ALDS) or ARC (ALDS related complex);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fluid or organs to neutralise/remove immunosuppressive peptides and/or viruses; or treating hepatitis A, B or C. The pharmaceutical compositions also treat a viral infection in a human or animal host. The human recombinant LDL receptor is also useful for manufacturing medicaments for reacting all the conditions given above. The human recombinant LDL receptor is a highly specific inhibitor of HIV-1 replication in vitro. (Updated on 17-0CT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enhancing the immune response using a recombinant human low-density lipoprotein receptor, useful for treating viral infections, especially human immunodeficiency virus (HIV) infection.
                                                                                                                                                 HIV; LDL; low density lipoprotein; human; immune response; infection; immunodeficiency; neoplastic tissue; myalgic encephalomyelitis; ME; viral infection fatigue syndrome; tuberculosis; hepatitis; AIDS; ARC; acquired immunodeficiency syndrome; AIDS related complex; HIV-infected CD4 cell; immunosuppressive peptide.
                                                                                                             HIV LDL binding peptide, sequence "A" variant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 12; 24pp; English.
                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-IB000149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-0072980P
                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PREN/) PRENDERGAST P T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-494040/41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prendergast PT;
                                                                                                                                                                                                                                                                                                                                            WO9938524-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                       28-JAN-1999;
                                           17-0CT-2003
19-0CT-1999
                                                                                                                                                                                                                                                                                                                                                                                         05-AUG-1999
AAY22583;
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This sequence represents a fragment of a HIV1 protein, and can be used in the immunogenic composition of the invention. The composition comprises a synthetic fusion polypeptide which includes a sequence encoding 1 or more T cell epitopes and a sequence encoding 1 or more B cell epitopes and a carrier. Both the T cell and B cell epitopes are derived from HIV proteins. The composition are useful as vaccines against HIV infection. The composition induces HIV-1-specific polyclonal antibodies that are opsonising and antiviral The peptide components may be selected induce a response against different viral isolates and in subjects who recognise different T cell epitopes. (Updated on 17-OCT-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Macrophage recruitment; chemokine derivative; MCP-1; osteoporosis; monocyte chemoattractant protein-1; inflammation; atherosclerosis; HIV; MIDS; stroke; psoriasis; autoimmune disease; hypertension; endotoxaemia; basophil-mediated disease; myocardial infarction; acute ischaemia; rheumatoid arthritis; contraception.
                                                                                                                                                                                                                                                                                                                                              100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 0.00014; Pred. o; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human chemokine derived peptide #25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB15873 standard; peptide; 24 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIQRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                    15; Conservative
                                                                                                                                                                                                                                                      standardise OS field)
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Best Local Similarity
                                                                                                                                                                                                                                                                                                      Sequence 24 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 77; DB 2; Length 24; 100.0%; Pred. No. 0.00014;
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Synthetic

AAY39769 standard, peptide; 24 AA.

RESULT 90 AAY39769 ID AAY3 XX

g 8

Local Similarity

Query Match Matches

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Gaps

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Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency virus, useful antigens in producing vaccines.
                                                      composition; T cell epitope; B cell epitope;
                                                                                                                                                                                                                                                       Disclosure; Col 73-74; 43pp; English.
                                                                              Human immunodeficiency virus 1.
                                                       HIV; vaccine; immunogenic comp
infection; antibody; antiviral
                                                                                                                                                                                      Sia CDY;
                                                                                                                                             93US-00073378.
                                                                                                                               95US-00467881
                                                                                                                                                                      (CONN-) CONNAUGHT LAB LTD.
                        (first entry)
                (revised)
                                       HIV1 chimeric peptide
                                                                                                                                                                                      Chong P,
                                                                                                                                                                                                       WPI; 1999-550482/46.
                                                                                                                               06-JUN-1995;
                                                                                                                                              09-JUN-1993;
                        26-NOV-1999
                17-OCT-2003
                                                                                                               14-SEP-1999.
                                                                                               JS5951986-A
                                                                                                                                                                                     Klein MH,
 AAY39769;
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Synthetic peptide(s) as vaccines for AIDS - selected from peptide regions which can fold as a maximally amphipathic helix recognised by I cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cornette JL, Ouyang CS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (USDC ) US SEC OF COMMERCE.
(USSH ) US DEPT HEALTH & HUMAN SERVICE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAP82464 standard; protein; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide component of AIDS vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               86US-00947935.
87US-00014430.
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    Zalipsky S, Woodle MC,
                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Delisi C, Margalit H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AIDS vaccine, T-cells.
                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                         WPI; 2001-201897/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NPI; 1988-184640/27
                                                                                                                                                                                                                                                                                                            Sequence 24 AA;
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12-FEB-1987;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP273716-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
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                                                                                                                                                                                    New peptide 3, amide and heterocyclic compounds and saccharide conjugates used for inhibiting chemokine induced activity and for treating e.g. stroke, vascular diseases, autoimmune diseases and tumor growth.
                                                                                                                                                                                                                                                           chemokines which can be used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences AABIS783-BIS794, AABIS803-BIS813 and AABIS831-BIS848. These chemokine derivatives can be useful or treat diseases such as autoimmune diseases, atherosclerosis, osteoporosis, HIV infection and diseases, endotoxaemia, myocardial infarction, acute ischaemia and rheumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in gene therapy for the same diseases, as well as in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                   The present invention concerns the identification of a number of
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                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 77; DB 3; Length 24; 100.0%; Pred. No. 0.00014; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV gp120 V3 loop; liposome composition; HIV infection.
                                                                                                                                                                                                                             Disclosure, Fig 18; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB68602 standard, peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus 1
                                                                         99US-00229071.
99US-00271192.
99US-00452406.
                                                    .2-JAN-2000; 2000WO-US000821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV gp120 V3 loop peptide #2.
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94US-00316436.
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                                                                                                                                          Tatalick LM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
                                                                                                                                                               WPI; 2000-499101/44.
                                                                                                                    (NEOR-) NEORX CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 AA;
           WO200042071-A2
                                                                                                                                                                                                                                                                                                                                                                        animal models
                                                                                                                                         Grainger DJ,
                                                                          12-JAN-1999;
                                                                                    17-MAR-1999;
01-DEC-1999;
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25-APR-2001
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                                20-JUL-2000
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29-SEP-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB68602;
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Gaps

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Mismatches

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Length 24; 0; Indels

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The present invention relates to a liposome composition comprising liposomes having an outer surface layer of polyethylene glycol chains, each having a free distal end. A polypeptide or polyeaccharide effector molecule is covalently attached to a portion of the distal ends. The effector interferes with specific binding of pathogen or cell in a bloodstream to a target cell or cell matrix, and is rapidly removed by renal clearance from the bloodstream, when administered in free form. The liposome composition may be used in treating a condition mediated by binding a pathogen or cell in the bloodstream, to a target cell or cell matrix. It can be used in treating septic shock, toxic shock, colonic inflammation, leukaemic cell proliferation, or HIV infection. The present sequence is a peptide of the V3 loop of HIV envelope protein gpl20. This captide may be used in the composition of the present invention. gpl20 binds to the CD4 receptor during HIV infection of lymphocytes. By introducing the present peptide, the CD4 receptors are blocked, thereby inhibiting HIV infection. (Updated on 11-SEP-2003 to standardise OS
                                                                                                                                                                                                                                       Liposome composition for use in treating septic shock comprises liposomes having an outer surface layer of polyethylene glycol chains, and a polypeptide or polysaccharide effector molecule.
Barenholz Y;
Martin FJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Fig 13; 32pp, English.
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AIDS; ARC; conjugate immunogen; Neisseria outer membrane protein;
HIV major neutralisation determinant.
                                                                                                                                                                                                              HIV peptide fragment (IIIB isolate)
                                                                              AAR08276 standard; protein; 25 AA.
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89US-00362177.
89US-00362178.
89US-00362179.
                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    90EP-00306082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Marburg S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1990-370100/50.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR13120;
                                                                                                                        AAR08276;
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                                      RESULT 95
                                                              AAR08276
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein derivative stimulates a lymphocyte proliferative response in HIV-
                                      This peptide is a component of an AIDS vaccine. It can fold as a maximally amphipathic helix and is recognised by T-cells immune to the AIDS virus envelope protein. See also AAP8462-63 and AAP82465-79. (Updated on 25-MAR-2003 to correct PA field.)
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                                                                                                                                                                                                                                      Gapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New HIV proteins and peptide(s) - used in diagnosis, prophylaxis or therapy of AIDS, esp. for prepn. of vaccines against HIV infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Revised record issued on 09-SEP-2004 : Correction to location
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 77; DB 1; Length 25; 100.0%; Pred. No. 0.00014; Ative 0; Mismatches 0; Indels
                                                                                                                                                                                     100.0%; Score 77; DB 1; Length 25; 100.0%; Pred. No. 0.00014; wative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV; AIDS; env gene; HIV vaccine; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Simian-Human immunodeficiency virus.
Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                  AAP90281 standard; protein; 25 AA
Claim 9; Page 10; 16pp; English
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                                                                                                                                                                                                                                                                              1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                       RIORGPGRAFVTICK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide 135 of HIV env gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
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                                                                                                                                                                                                              Best Local Similarity 100.
Matches 15; Conservative
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Petro J, Okeeffe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
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                                                                                                                                                 Sequence 25 AA;
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22-JUN-1990
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AAP90281

AAP90281

DT 25-P

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EP90 OS Simion OS S
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Larson VM;

Scolnick EM,

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composed of neutralising
                                                                                               This peptide is derived from the HIV IIIB isolate and is cross-reactive with the HIV major neutralisation determinant (MNLD). This MNLD is used in a conjugate, covalently linked to the outer membrane protein (Omp) from Neisseria, as an immunogen for vaccination against AIDS. A cocktail of different MNLD poly- peptides can be used. See also AAR08274-75 and AAR08277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-idiotypic; antibody; gp120; HIV; human immunodeficiency virus; paratope; complemetarity determining region; CDR; immunisation; vaccine; immunotoxin; T-cell; AIDS; ARC.
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                                                                                                                                                                                                                                                                                                          100.0%; Score 77; DB 2; Length 25; 100.0%; Pred. No. 0.00014;
                                                                                                                                                                                                                                                                                                                                                   0; Indels
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Conjugate immunogen for AIDS and ARC treatment -determinant of HIV and Neisseria outer membrane.
                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR13120 standard; peptide; 25 AA.
                                                                    Claim 2; Page 22; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIORGPGRAFVTIGK 15
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1 RIQRGPGRAFVTIGK 15 

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This peptide is one example of an HIV-1 amplifier peptide for use in a composition for enhancing the immunogenicity of an envelope glycoprotein of a virus. The sequence corresponds to the major neutralisation epitope (loop V3) of HIV-1 brull isolate and enhances the induction of persistent neutralising antibodies in the host. The amplifier peptide is need in addition to an envelope glycoprotein for priming the induction of neutralising antibodies. The compositions are particularly useful for vaccinating against HIV, SIV, HILV-I and HILV-II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a co-conjugate comprising an immunogenic protein or protein complex having a first set of covalent linkages to low molecular weight moieties which have an anionic or polyanionic character at physiological ph, and a second set of covalent linkages to peptides comprising HIV principal neutralizing determinants (NND's) or immunologically equivalent peptides. Preferably at least one set of the covalent linkages is comprised of maleinide derivatives; the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "bonds to the OMPC of the conjugate via this site"
  for use as vaccine
Enhancing immunogenicity of envelope glyco:protein - for use as vaccin
or immuno:therapeutic drug especially against HIV, HTLV-I and HTLV-II.
                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                meningitidis b; outer membrane protein complex; OMPC; PND135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human immunodeficiency virus; vaccine; Neisseria;
                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 77; DB 2; Length 25; Best Local Similarity 100.0%; Pred. No. 0.00014; Matches 15; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR31276 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV principal determinant peptide
                                                            Claim 13; Page 50; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 12; Page 56; 63pp; English.
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90US-00555966.
91US-00715276.
91US-00715278.
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preparing vaccines
                                                                                                                                                                                                                                                                                   Sequence 25 AA;
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Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AIDS; ARC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                     The peptide corresponds to residues 294-318 of the gpl20 envelope protein of HIV-1 which is a principal neutralising determinant (PND). Abs recognise residues 294-308 (MAb BAT267) or 304-318 (MAb 125). These MAb are used to raise anti-idiotypic Abs (AAbs). The Abs are useful for passive immunisation and as components for immunotoxins which destroy T-cells infected with HIV. They inhibit T-cell infection and syncytium formation, are group specific and neutralise specific strains of HIV-1. They can be used to treat AIDS or ARC. The AAbs can be used for active immunisation or can be admin with another vaccine to increase antigenicity. See also AAR13121. (Updated on 24-OCT-2003 to standardise
                                                                                                                                                                                                                                                                                                                       Monoclonal antibodies specific to the gp120 HIV envelope protein - for immunisation against HIV in treatment of AIDS or ARC.
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                                                                                                                                                                                                                                                                                                                                                                               Claim 5; Page 97; 124pp; English
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                                                                                                  89US-00454161.
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90US-00531789
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                                                                                                                                                                                                                                          Chang TW, Fung MSC,
                                                                                                                                                                                                                                                                                 WPI; 1991-222664/30.
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les 15; Conser
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                                                                                                  21-DEC-1989;
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12-JUN-1990;
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Matches

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RESULT 97 AAR15058

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Gaps

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The peptide is HIV principle neutralising determinant (PND) 135 and is used as part of a conjugate comprising the major immune enhancing protein (MIEP) of Neisseria meningitidis covalently linked to the HIV PND. The conjugate may be used to prepare vaccines against HIV infections, e.g. AIDS, as research tools for studying PND structure- function relationships, or as immunogens for use in the passive immunisation of humans. (Updated on 25-MAR-2003 to correct PN field.)
            form of a carboxylic, sulphonic or phosphonic acid; the immunogenic protein is the outer membrane protein complex (OMPC) of Noisseria meningitidis b; and the PND peptide has a linear structure, a disulphide-bonded cyclic structure. The present sequence (PND135) is an example of a bonded cyclic structure. The present sequence (PND135) is an example of a byd peptide component used in the co-conjugate. The co-conjugate is useful for inducing anti-peptide immune response in mammals, for inducing prevent HIV infection or disease, including AIDS, or for treating humans afflicted with HIV infection or disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus; AIDS; PND; MIEP; conjugate; major immune enhancing protein; vaccine; anti-HIV antibodies; immunogen; passive immunisation.
composed of one to five residues of the anionic
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                                                                                                                                                                                                                 100.0%; Score 77; DB 2; Length 25; 100.0%; Pred. No. 0.00014;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus.
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                                                                                                                                                                                                                                                  15; Conservative
   (poly) anionic moiety is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MERI ) MERCK & CO INC.
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                                                                                                                                                                                                                                  Best Local Similarity
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28-APR-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV peptide-polysaccharide-protein conjugates - used in vaccines or to produce antibodies to prevent or treat HIV infection.

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    /note= "Joins onto polysaccharide-protein complex via

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                                                                                                                                                                                                                       Human immunodeficiency virus; principal neutralizing determinant;
outer membrane protein complex; OMPC; Neisseria; AIDS; PND135.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 77; DB 2; L 100.0%; Pred. No. 0.00014; tive 0; Mismatches 0;
                                                                                                                                                                                            HIV-PND-polysaccharide-protein conjugate vaccine.
                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                     AAR26712 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Emini EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 9; Page 57; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90US-0055558.
90US-00555974.
91US-00715275.
91US-00715277.
                                                                                                                                                                                                                                                                                                                                                                                                                                            90US-0055558
1 RIORGPGRAFVTIGK 15
                          8 RIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8 RIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                 this site"
                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Marburg S, Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1992-034437/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25 AA;
                                                                                                                                                                                                                                                                                                     Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-1990;
                                                                                                                                                                 09-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .9-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                              29-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .9-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                 EP468714-A.
                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                   AAR26712;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Loca
Matches
                                                                        RESULT 100
                                                                                       AAR2671
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100.0%; Score 77; DB 2; Length 25; 100.0%; Pred. No. 0.00014; ative 0; Mismatches 0; Indel8

Query Match Best Local Similarity 100. Matches 15; Conservative

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Disclosure; Page 3; 35pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR41330 standard; peptide; 25 AA.
                            Human immunodeficiency virus 1.
                                                                                                      92JP-00098602.
92JP-00237648.
93JP-00054239.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93WO-US002349.
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                                                                                    93WO-JP000327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                             1 RIORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100...
Pest Local Si Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WOHL/) WOHLSTADTER J N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
                                                                                                                                                                                   WPI; 1993-320455/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV gp120 epitope.
                                                                                                                                                                                                                                                                                                                                                    Sequence 25 AA;
                                                                                                                                             (TSDT-) TSD KK
                                              WO9318791-A1
                                                                                    19-MAR-1993;
                                                                                                                          15-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     409319170-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
21-APR-1994
                                                                  30-SEP-1993
                                                                                                                                                                                                                                                                                                                                 OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR41330;
                                                                                                                                                                 Okuda K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR41330
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                                                                                                                                                                                                                                                                                                                                                                                                  The sequence is that of peptide RP135 (IIIB) used as an immunogen for the generation of antibodies directed against the V3 loop of HIV gp120. These antibodies can be used as part of a compsn. With antibodies directed against the CD4 binding site of gp120. The antibodies act synergistically to neutralise HV-1 in the treatment of HIV infection caused by different strains. (Updated on 25-MRR-2003 to correct PN field.) (Updated on 25-MRR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                          Synergistic compsn. for treating HIV-1 infection - comprises antibody to V3 loop of GP120 and antibody to CD4 binding site of GP120 or soluble CD4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         V3 region; HIV; envelope gp120; vaccine; human; humoral response;
                                                                                                       HIV-1; human immunodeficiency virus; antibody generation; AIDS; infection; CD4 binding site; soluble CD4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                 25
/note= "not in natural sequence of isolate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 77; DB 2; Length 25; 100.0%; Pred. No. 0.00014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IndelB
                                                                                    HIV gp120 V3 loop immunogenic peptide RP135 (IIIB).
                                                                                                                                                                                                                                                                                                      Herlihy WC;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                      Field KG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV gp120 V3 region peptide HIV-III B.
                                                                                                                                                      Location/Qualifiers
                  AAR33222 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR41336 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                 Example; Page 12; 56pp; English.
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                                                                                                                                                                                                                                                     91US-00756677.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22
                                                                                                                                                                                                                                                                                                      Potts BJ, Whitescharf ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8 RIORGPGRAFVTIGK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
                                                                                                                                                                                                                                                                                 (REPK ) REPLIGEN CORP
                                                                                                                                                                                                                                                                                                                       WPI; 1993-100653/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25 AA;
                                                                                                                                                                                                                                                      09-SEP-1991;
20-JUL-1992;
                                                                                                                                                                                                                                   02-SEP-1992;
                                                                                                                                                                                             WO9304693-A1
                                                                                                                                                                                                                                                                                                                                                              polypeptide.
                                                        25-MAR-2003
13-JUL-1993
                                                                                                                                                                                                                18-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
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                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR41336;
                                     AAR33222;
                                                                                                                                                                 Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 102
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Virus for prevention of HIV infected diseases - comprising several peptide(s) consisting of V3 region peptide of envelope Gp., 120, etc. and complex including carrier protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        derived from the V3 region of HIV envelope gp120. These peptides may be used in a vaccine which is effective in humans and animals and activates humoral and cellular immunity. The vaccine also contains a carrier protein containing a cysteine group, eg. human serum albumin (HSA), keyhole limpet haemocyanin (KLH) or multiple antigen peptide. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 24-OCT-2003 to standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequences given in AAR41336-39 and AAR42664 represent peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV; haemagglutinin; reactants; catalysts; cofactors; repressors; enhancers; hormones; binders; human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
cellular immunity; carrier protein; human serum albumin; HSA;
keyhole limpet haemocyanin; KLH; multiple antigen peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 77; DB 2; Length 25; 100.0%; Pred. No. 0.00014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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the virus and to confer long lasting immunity, longer than 6 m also AAR36567-86. (Updated on 25-MAR-2003 to correct PN field.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW72819
   SSXS
                                                                                                                                                                                                                                        The HIV gpl20 epitope is used to isolate, create or evolve novel mols. including (in) organic and biomolecules such as proteins, peptides, nucleic acids, oligonuclectides, lipids, and polysaccharides for use as reactants, catalysts, enzymatic cofactors, repressors, enhancers, hormones and binders for a wide variety of substrates in industrial and therapeutic products. This epitope was isolated from variable region 3 of HIV gpl20 (amino acids 271-295). (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ô
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administration of viral envelope glycoprotein itself, and an oligopeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 glycoprotein (EGP) of a virue (esp. HIV gp120 or gp160) in a host comprises admin. to the host at least one EGP of the virus in an amt. sufficient for priming vaccination and at least one peptide derived from an amino acid sequence of the EGP (e.g. the sequence shown), where the peptide comprises at least one virus-neutralisation epicope (VNE). The complex is able to enhance the induction of neutralising antibodies to
                                                                                                                        Obtaining a novel mol. - capable of a desired interaction with a substrate of interest and a selection molecule expressed by the host.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus; gp120; gp160; EGP; VNE; immunity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Virus neutralising epitope of envelope glycoprotein of HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 77; DB 2; Length 25; 100.0%; Pred. No. 0.00014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 82; 107pp; English.
                                                                                                                                                                                                                  Claim 151; Page 147; 165pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR36587 standard; peptide; 25 AA.
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                                                             WPI; 1993-320737/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 25 AA;
         Wohlstadter JN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
06-SEP-1993
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28-OCT-1991;
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The present invention describes an antibody conjugate comprising an antibody (Ab) which binds to an epitope within amino acid residue 308-322 of gpl20 and neutralises HIV-1, conjugated with a cytotoxic agent, an anti-viral agent or an agent which facilitates passage through the blood brain barrier. Also described is an antibody conjugate as above but where the Ab binds to an epitope within amino acid residue 298-312 of gpl2 which neutralises HIV-1. The present sequence represents an HIV-1 gpl20 epitope corresponding to positions 294 to 318. The Ab are monoclonal Ab which bind to the gpl20 protesin on the envelope of HIV-1. They inhibit the infection of T-cells and also inhibit syncytium formation. The cantibodies are group specific and neutralise different scrains and isolates of HIV-1. The antibodies have a variety of uses, including the treatment and prevention of AIDS and AIDS related complex. They are especially used to kill infected T-cells. (Updated on 17-OCT-2003 to
                                                             ö
                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antibody conjugate comprising monoclonal antibody - which binds to epitope within amino acid residue of gp120 which neutralises HIV-1 conjugated with, e.g. cytotoxic agent.
                                                                                                                                                                                                                                                                                                                                                         HIV-1; gpl20; epitope; monoclonal antibody; envelope; neutralise; inhibit; infection; T-cell; inhibit syncytium formation; AlDS.
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                            100.0%; Score 77; DB 2; Length 25; 100.0%; Pred. No. 0.00014;
                                                             0; Indels
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                                                             0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                             AAW72819 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          1. .15
/label= peptide_a
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/label= peptide_b
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                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                              HIV-1 gp120 epitope 294 to 318.
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87US-00137861.
89US-00343540.
92US-00895197.
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                                                                                             1 RIQRGPGRAFVTIGK 15
                                                                                                                RIORGPGRAFVTIGK 22
                                                                                                                                                                                                                                                                                             (first entry)
                                                             15; Conservative
                                                                                                                                                                                                                                                                              (revised)
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                                               Similarity
Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAY-1987;
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13-JAN-1999
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                            Query Match
Best Local S:
Matches 15
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Indels

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Pred. No. 0.00014; Mismatches 0;

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11 RIORGPGRAFVTIGK

100.0%;

Best Local Similarity 100. Matches 15; Conservative

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents an epitope of the gp120 protein of human immune deficiency virus (HIV)-1. The sequence comprises amino acids 298 to 322 of gp120. The specification describes monoclonal antibodies which bind to sequences derived from the present epitope. Specifically, these antibodies are designated BAT123, 267 and 085. Monoclonal antibodies are designated BAT123, 267 and 085. Monoclonal antibodies ownernalise HIV-1, inhibiting both infection of T cells and formation of syncytia, so are used to treat acquired immune deficiency syndrome (AIDS) and AIDS-related complex, by passive immunisation, as carriers of cytotoxic or antiviral agents, and in extracorporeal systems. They can also be used as immunoassay reagents (for diagnosis or measurement of viral load) and to screen for neutralising epitopes, potentially useful in vaccine development. (Updated on 20-MAR-2003 to correct PR field.) (Updated on 17-0CT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monoclonal antibodies directed against regions of gp120 of human immune deficiency virus-1 - are neutralising and able to inhibit infection of ocl1s and formation of syncytia, used for treatment, prevention or diagnosis of acquired immune deficiency syndrome.
                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                   Epitope; gpl20 protein; monoclonal antibody; HIV-1; antibody BAT123; antibody BAT267; antibody BAT1085; T cell infection inhibition; syncytia formation; acquired immune deficiency syndrome; AIDS; AIDS-related complex; passive immunisation; antiviral; cytocoxic;
                                                                                                         ò
                                                                   100.0%; Score 77; DB 2; Length 25; 100.0%; Pred. No. 0.00014; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chang TW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chang NT,
                                                                                                                                                                                                                                                            AAW87618 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         viral load measurement; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                  Epitope of HIV-1 gp120 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87US-00137445.
87US-00137861.
91US-00767533.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sun CRY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Col 8; 16pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92US-00950571
                                                                                                                                      1 RIORGPGRAFVTIGK 15
                                                                                                                                                                        11 RIORGPGRAFVTIGK 25
                                                Query Match
Best Local Similarity 100...
Best Local Similarity 100...
                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                 (revised)
standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fung MSC, Sun BNC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-095002/08
                                Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (TANO-) TANOX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-DEC-1987;
26-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-MAY-1987;
                                                                                                                                                                                                                                                                                                                                 17-OCT-2003
                                                                                                                                                                                                                                                                                                                                               20-MAR-2003
03-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US5854400-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-DEC-1998
                                                                                                                                                                                                                                                                                            AAW87618;
                                                                                                                                                                                                                           RESULT 106
AAW87618
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100.0%; Score 77; DB 2; Length 25;

Sequence 25 AA;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunostimulant peptide, used as an anti-carcinoma vaccine, comprises a an epitope of the non-VNTR, non-leader region of a mucin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                       Mucin, cytostatic, immunostimulant, cell mediated immune response; carcinoma; adenocarcinoma; breast cancer; dendritic cell; vaccine; gene therapy, CTL; cytotoxin T-lymphocyte.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 77; DB 4; Length 25; 100.0%; Pred. No. 0.00014; ive 0; Mismatches 0; Indels
                                                                                                                                   Human immunodeficiency virus Dd haplotype peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Apostolopoulous V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 19; 84pp; English
                                AAE09522 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR04427 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                 01-FEB-2000; 2000AU-00005369.
14-JUN-2000; 2000US-00593870.
                                                                                                                                                                                                                                                                                                                                                01-FEB-2001; 2001WO-AU000090.
                                                                                                                                                                                                                                          Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ackenzie IFC, Pietersz GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 RIORGPGRAFVTIGK 19
                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-541537/60
                                                                                                                                                                                                                                                                           WO200157068-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25 AA;
                                                                                                    19-NOV-2001
                                                                                                                                                                                                                                                                                                               09-AUG-2001.
                                                                   AAE09522;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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RESULT 107
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                  AAE09522
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                                                  요
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us-08-869-386-1.rag

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Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAP). In peptide 18-4, the Arg residue at position 4 in peptide 18 has been replaced by a Lys residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                         Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                                                                                                                                                                                Shirai M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96.1%; Score 74; DB 2; Length 15; 93.3%; Pred. No. 0.00025;
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                                                                                                                                                                                                                                                                                                                                Nara P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Mismatches
                                                                                                                                                                                                                                                                                                                                Pendleton CD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 33; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR66430 standard; peptide; 15 AA.
                                                                                                                                                                                                        94WO-US005142.
                                                                                                                                                                                                                                               93US-00060988.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                        (USSH ) US SEC DEPT HEALTH.
                                                                                                                                                                                                                                                                                                                                Berzofsky JA, Ahlers JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                       WPI; 1995-006707/01.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 AA;
                                                                                                                                                                                                                                                 14-MAY-1993;
                                                                                                                                                                                                        13-MAY-1994;
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                                                                                                                         WO9426785-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-AUG-1995
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                                                                                   Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 110
AAR66430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
    %XXCCCCCCCCCXXXX474474X6X6X6X6X6X6X6X6X6X8X8X8X8X8X8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptide 135 comprises segments of the Principal Neutralising Domain (envelope protein) from isolate HIV-IIIB. The last Cys resisue is added for the purpose of crosslinking to carrier proteins. Cysteine residues can be added so that that residues are nor near both ends form a disulfide bond, thus giving the peptide a loop-like configuration, which is utilised to enhance the immunogenic properties of the peptide. The peptide is capable of eliciting, and/or binding with, neutralising antibodies. The neutralising domain is bounded by cysteine residues which occur at positions 296 and 331. Peptides can be used as immunogens or screening reagents to generate or identify poly- or MAbs. See also AAR04427-R04506 and AAR04277- (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Revised record issued on 09-SEP-2004 : Correction to Peature Table Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Principal neutralising domain of HIV variants - used for producing peptide(s) and antibodies for diagnosis, prophylaxis and/or therapy
                                                                                                                                          HIV-IIIB; peptide 135; principal neutralising domain; antibodies;
diagnosis; prophylaxis; therapy; AIDS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Grimaila R;
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93.3%; Pred. No. 0.00028;
ive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Javaherian K, Farley J,
                                                                                                    Human immunodeficiency virus peptide 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 8 (30); Page 75; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR66419 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy therapy of HIV infection.
                                                                                                                                                                                                                                                                                                                                                                     88US-00252949.
89US-00359543.
89US-00407663.
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                       (revised)
(revised)
(first entry)
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(first entry)
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Matches 14; Conservative
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    Putney SD,
Petrobre J;

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                     09-SEP-2004
25-MAR-2003
20-SEP-1990
                                                                                                                                                                                                                                                                                                                                03-OCT-1988;
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                                                                                                                                                                                                                                                 WO9003984-A
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03-AUG-1995
                                                                                                                                                                                                                                                                                          19-APR-1990
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                                                                                                                                                                                                           Synthetic.
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Query Match

Lynn DU,

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Gape

Shirai M;

Nara P,

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In the HIV-1 IIIB sequence (AARG6414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCIOS 3-18 and PCIOS 6-18 (see AARG6419). Analysis of the substituted peptides (AARG6416-86649) showed that binding was enhanced over peptide 18 control when a tyrosine was substed. For a Val at position 11 and substems at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising position 15 in peptide 18 has been replaced by a GIN residue at 25-WAR-2003 to correct PN field.)
                                                                                                                    Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                            Single residues from the HIV-1 RF sequence (AAR66415) replaced residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                    94.8%; Sco...
93.3%; Pred. No. v...
1; Mismatches
                                                              Pendleton CD,
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93US-00060988
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                            (USSH ) US SEC DEPT HEALTH
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                                                           Berzofsky JA, Ahlers JD,
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                                                                                         WPI; 1995-006707/01
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 AA;
14-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-MAY-1993;
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Nara P,

Gaps ö

Length 15; 0; Indels

Score 73; DB 2; Pred. No. 0.00036;

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in the HIV-1 IIIB sequence (ARR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCUS 3-18 and PCUUS 6-18 (see ARR66409 and AAR66411). Analysis of the substituted peptides (AAR66416- R66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a val at position 11 and substras. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of determinant sequence (PGRAP). In peptide 18-9, the Ala residue at position 9 in peptide 18 has been replaced by a Val residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a polyfunctional base sequence (N1) avaing two or more functions in different reading frames. Also described are: (1) a method for producing N1 and artificial gene expression vectors comprising N1; (2) transgenic non-human animals comprising N1; and (3) artificial comprising N1; and (3) artificial tissues or high molecular weight artificial protein, artificial to creating industrially useful artificial matrix proteins for each culture. The present sequence represents a peptide which is used in an example from the present invention. (Updated on 29-AUG-2003 to standardise OS field)
                                 Single residues from the HIV-1 RF sequence (AAR66415) replaced residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polyfunctional base sequence having two or more functions in different reading frames, useful for producing artificial matrix proteins for cell
                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polyfunctional base sequence; microgene; industrial; cell culture; artificial matrix protein; transgenic animal; HIV.
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                                                                                                                                                                                                                                                                                                          Length 15;
                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                        Score 73; DB 2; I
Pred. No. 0.00036;
                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV gp120 related peptide SEQ ID NO:1.
Example 1; Page 33; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NISC-) JAPAN SCI & TECHNOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 46; 61pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB05775 standard; peptide; 20 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JUN-2001; 2001WO-JP005116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-JUN-2000; 2000JP-00180997
                                                                                                                                                                                                                                                                                                                                                                                                  1 RIQRGPGRVFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                            1 RIQRGPGRAFVTIGK 15
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(first entry)
                                                                                                                                                                                                                                                                                                                                          14; Conservative
                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                      Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200196558-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-AUG-2003
07-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABB05775;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   culture.
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(first entry)
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                                                                                                                              13-MAY-1994;
                                                                                                                                                 14-MAY-1993;
03-AUG-1995
                                                                                          WO9426785-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-AUG-2003
30-MAR-1992
                                                                                                            24-NOV-1994
                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                  PN field.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HTLV-IIIB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAP95357;
                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                             The invention comprises an artificial protein which induces a strong timmune response to a peptide group (the protein contains all or part of the peptide group). The artificial protein assists the formation of high-order protein structure and/or assists the antigen presentation of immunocompetent cells. The artificial protein of the invention is useful for inducing a strong immune response and the preparation of effective antibodies to specific antigens, especially HIV. The present amino acid sequence represents a peptide that was used in the invention
                                                                                                                                                                                              Strong immune response induction; high-order protein structure formation;
                                                                                                                                                                                                                                                                                                                                                                        Artificial protein capable of inducing a strong immune response to a peptide group for assisting antibody production in vivo to viruses and other antigens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                    Gaps
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        Score 73; DB 5; Length 2v, Pred. No. 0.00047;
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                                                                                                                                                                            Strong immune response induction-related peptide 1.
                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                    (NISC-) JAPAN SCI & TECHNOLOGY CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR66416 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94.8%; Scor.
93.3%; Pred
0; 1
                                                                                                                      AA015657 standard; peptide; 20 AA
                                                                                                                                                                                                                                                                                                                                                                                                               Claim 6; Page 5; 77pp; Japanese.
                94.8%;
                                                                                                                                                                                                                                                                                                  13-OCT-2000; 2000JP-00314288
                                                                                                                                                                                                                                                                                10-OCT-2001; 2001WO-JP008893
                                                                        RIORGPGRTFVTIGK 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RIORGPGRTFVTIGK 19
                                                       1 RIQRGPGRAFVTIGK 15
                                                                                                                                                           08-NOV-2002 (first entry)
                                                                                                                                                                                                       antigen presentation; HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 93.3
hes 14; Conservative
                                    14; Conservative
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                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 AA;
Sequence 20 AA;
                                                                                                                                                                                                                                            WO200233074-A1
                                                                                                                                                                                                                          Unidentified
                                                                                                                                                                                                                                                              25-APR-2002
                                                                                                                                         AA015657;
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                                                                                                                                                                                                                                                                                                                                      Shiba K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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AAR66416
ID AAR664
XX
AC AAR664
XX
DT 25-MAF
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Matches
                                    Matches
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in the HIV-I IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals munuised with the cluster peptides PCLUS 3.18 and PCLUS 6.18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-R66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-1, the N-terminal residue that in peptide 18 has been deleted. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Variable region V3, found in the envelope protein {	t gp120} of an AIDS or ARC causing or related virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Single residues from the HIV-1 RF sequence (AAR66415) replaced residues
                                                               T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps

    .15
/note= "an example of a peptide of the invention"

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shirai M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Д,
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100.0%; Pred. No. 0.00048;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Berzofsky JA, Ahlers JD, Pendleton CD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 33; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vaccine; AIDS; ARC; HIV; diagnosis
HIV-1 IIIB peptide 18-1 (316-330).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAP95357 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                        94WO-US005142.
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(first entry)
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Matches 14; Conservative
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Disclosure, Page 62; 106pp; English
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                                               WPI; 1993-093577/11
                                                                                                                                                                                                                          Sequence 15 AA;
                                                                                    glyco:protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-MAR-1994;
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03-MAY-1995
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                                                                                                                                                                                                                                                                                                                                                                AAR62166;
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Best Local S
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                                                                                                                                                                                                                                       The peptides of the invention comprise the beta-turn AA SQ GPG or GPGR at positions 312-314 or 312-315 in the AA numbering of HTLV-IIIB (BH10) and flanking AA SQs having a length equal to or greater than 1 and pref. equal to or greater than 2 AAs; variants in which the GPG or GPGR SQ has been replaced by a different beta-turn SQ; and variants in which the free NH2-terminal gp AA and/or the free carboxyl terminal gp. AA has been blocked or modified otherwise. (Updated on 27-AUG-2003 to correct OS
                                                                                                                                                                                           Oligopeptide(s) corresp. to beta-turn variable region of gp.120 - used for diagnosis of and prodn of vaccines against AIDS and ARC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence of synthetic peptide which represents residues 315-329 of the pg160 envelope protein of HIV-1 isolate IIIb.
                                                                                                                                                                                                                                                                                                                                                                 Gaps
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glycoprotein 160.
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                                                                                                                                                                                                                                                                                                                                            93.5%; Score 72; DB 1; Length 15; 100.0%; Pred, No. 0.00051; Ative 0; Mismatches 0; Indels.
         /note= "see above"
                  3. .12
/note= "see above"
                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR33460 standard; peptide; 15 AA
                                                                                                                                                                                                                       Disclosure; Page 3; 7pp; English.
                                                                                                                   (DIER-) STICHTING CENT DIER.
(UNAM ) UNIV VAN AMSTERDAM.
(UYAM-) UNIV AMSTERDAM ZIEKENHUI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus 1
                                                                                  88EP-00202248
                                                                                                   87NL-00002403
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                                                                                                                                                                                                                                                                                                                                                               14; Conservative
                                                                                                                                                          Goudsmit J, Meloen RH;
Misc-difference 3. .13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(revised)
(revised)
                                                                                                                                                                           WPI; 1989-108193/15
                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                 Misc-difference 3
                                                                                                                                                                                                                                                                                                                          Sequence 15 AA;
                                                                                 07-OCT-1988;
                                                                                                  09-OCT-1987;
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18-SEP-1991;
                                                              12-APR-1989
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25-MAR-2003
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                                             EP311219-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR33460;
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AAR33460
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                                                                           s;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                      Peptide(s) for stimulation of cytotoxic T cells specific for HIV-1 which correspond to residues 318-327 of HIV-1 gp 160 envelope
                                                                           Kozlowski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    epitope, autoantibody; immunoinfective cluster virus;
nuclear protein antigen; systemic rheumaciic disorder;
human immunodeficiency virus; HIV-1; systemic lupus erythematosus;
mixed connective tissue disease; scleroderma; glycoprotein 120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV-1 gp120 V3 loop dominant neutralising domain in chimpanzees.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ch 93.5%; Score 72; DB 2; Length 15; Similarity 93.3%; Pred. No. 0.00051; 14; Conservative 0; Mismatches 1; Indels
                                                                           Pendleton CD,
                                                                       Berzofsky JA, Taskeshita T, Shirai M,
(USSH ) US DEPT HEALTH & HUMAN SERVICE
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                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 9; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR62166 standard; peptide; 15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94WO-US002631.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
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Previous immunological analyses of the V3 loop of HIV-1 (AAR62159) have localised the main neutralising domains. The target of more than 80% of neutralising antibodies in HIV-1 infected sera from AIDS patients has now been found to overly the consensus binding sequence and domain A epitopes of the U1 shRNP 70% protein. In AIDS, antibody titres are too low to arrest the disease, however, the homologous sequences in 70% are immunodominant targets of autoantibodies in the systemic rheumatoid disorder of mixed connective tissue disease. The titers of such autoantibodies exceed 10 power 7. The anti-shRNP autoantibodies will cross-react with HIV-1 epitopes and are useful for treating HIV AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single residues from the HIV-1 RP sequence (AAR66415) replaced residues in the HIV-1 IIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-R64300) showed that binding was enhanced over peptide is control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of seara was reduced when substns. were made in the principal neutralising determinant sequence (PGRAP). In peptide 18-12, the Thr residue at position 12 in peptide 18 has been replaced by an Ala residue. (Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different WHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shirai M;
                                                                                                                                                                                                                                                                                                    93.5%; Score 72; DB 2; Length 15; 100.0%; Pred. No. 0.00051; Attive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nara P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ahlers JD, Pendleton CD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 33; 120pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR66427 standard; peptide; 15
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                                                                                                                                                                                                                                                                                                                                                                                                          2 RIORGPGRAFVTIG 15
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                                                                                                                                                                                                                                                                                                                                                                                    1 RIQRGPGRAFVTIG 14
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                                                                                                                                                                                                                                                                                                                                              14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV-1 IIIB peptide 18-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
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                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                    Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Berzofsky JA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
03-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                     T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
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                                                         Length 15
                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Berzofsky JA, Ahlers JD, Pendleton CD, Nara P,
                                                        Score 72; DB 2; 1 Pred. No. 0.00051; 0; Mismatches 1
on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 33; 120pp; English
                                                                                                                                                                                                                              AAR66428 standard; peptide; 15 AA.
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                                                         93.5%;
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                                                                                                                                                  1 RIORGPGRAFVAIGK 15
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                                                                                                                                                                                                                                                                                           (revised)
(first entry)
                                                                                                                     1 RIORGPGRAFVTIGK
                                                                                                                                                                                                                                                                                                                                          HIV-1 IIIB peptide 18-13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14; Conservative
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                            Sequence 15 AA;
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03-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                            AAR66428;
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Best Local &
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                                                                                                                                                                                                                  AAR66428
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Vaccine; AIDS; ARC; HIV; diagnosis.
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100.0%; Prev
0; F
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(UNAM ) UNIV VAN AMSTERDAM.
(UYAM-) UNIV AMSTERDAM ZIEKENHUI.
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                                                                                                92WO-US007511.
                                                                                                                     91US-00756677.
92US-00916542.
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                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RIQRGPGRAFVTIG
                                                                                                                                                    (REPK ) REPLIGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                              WPI; 1993-100653/12.
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 16 AA;
                                                    WO9304693-A1
                                                                                                                                20-JUL-1992;
                                                                                               02-SEP-1992;
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30-MAR-1992
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                                                                          18-MAR-1993
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                                Synthetic.
                                                                                                                                                                          Potts BJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HTLV-IIIB.
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                                                                                                                                                                                                                                                                                                                                                                                                               Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-AAR66409 showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of determinant sequence (PGRAP). In peptide 18-11, the Val residue at position 11 in peptide 18 has been replaced by a Tyr residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                             T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                                                                                                                                                        Berzofsky JA, Ahlers JD, Pendleton CD, Nara P, Shirai M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , DB 2; Lens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-IIIB gp120 V3 loop epitope peptide RP135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ed. No. 0.00
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 72;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 33; 120pp; English
                            AAR66426 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR33236 standard; peptide; 16 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93.5%;
93.3%;
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                                                                                   (first entry)
                                                                                                                                                                                                                                                                                  (USSH ) US SEC DEPT HEALTH
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(first entry)
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Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                        HIV-1 IIIB peptide 18-11.
                                                                         (revised)
                                                                                                                                                                                                                                                                                                                             WPI; 1995-006707/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 AA;
                                                                                                                            cell helper
                                                                                                                                                                                                                                         13-MAY-1994;
                                                                       25-MAR-2003
03-AUG-1995
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13-JUL-1993
                                                                                                                                                                       Synthetic.
                                                  AAR66426;
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      RESULT 120
AAR66426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 121
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ID AAR3

XX AC AAR3

XX DT 25-r

DT 13-c

DT 13-c

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The sequence is that of the HIV-IIIB V3 loop epitope peptide RP135 which and used in a competition assay to determine whether a given anti-V3 loop antibody will have strong neutralisation activity by itself, and if it has potential to act synergistically with a second agent. The assay can be used to test for potential neutralisation activity of any anti-V3 loop antibody towards any isolate by using a peptide defined from the V3 loop from the HIV isolate of interest as the competitor (Updated on 25-MAR-2003 to correct PI field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synergistic compsn. for treating HIV-1 infection - comprises antibody to V3 loop of GP120 and antibody to CD4 binding site of GP120 or soluble CD4 polypeptide.
HIV-1; human immunodeficiency virus; competition assay; AIDS; infection;
CD4 binding site; soluble CD4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Variable region V3 sequence found in the envelope protein gp120 of an ANDS or ARC causing or related virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 72; DB 2; Le; Pred. No. 0.00054; 0; Mismatches 0;
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The peptides of the invention comprise the beta-turn AA SQ GPG or GPGR at positions 312-314 or 312-315 in the AA numbering of HTLV-IIIB (BH10) and flanking AA SQs having a length equal to or greater than 1 and pref. equal to or greater than 1 and pref. been replaced by a different beta-turn SQ; and variants in which the GPG or GPGR SQ has been replaced by a different beta-turn SQ; and variants in which the free NH2-terminal gp AA and/or the free carboxyl terminal gp. AA has been
                                                                                                                                                   The peptides of the invention comprise the beta-turn AA SQ GPG or GPGR at positions 312-314 or 312-315 in the AA numbering of HTLV-IIIB (BH10) and flanking AA SQs having a length equal to or greater than 1 and pref. equal to or greater than 2 AAs; variants in which the GPG or GPGR SQ has been replaced by a different beta-turn SQ; and variants in which the free NH2-terminal gp AA and/or the free carboxyl terminal gp. AA has been blocked or modified otherwise. (Updated on 27-AUG-2003 to correct OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variable region V3 found in the envelope protein gp120 of an AIDS or ARC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligopeptide(s) corresp. to beta-turn variable region of gp.120 - used for diagnosis of and prodn of vaccines against AIDS and ARC.
                                                                          Oligopeptide(s) corresp. to beta-turn variable region of gp.120 for diagnosis of and prodn of vaccines against AIDS and ARC.
                                                                                                                                                                                                                                                                                                                                 93.5%; Score 72; DB 1; Length 17; 100.0%; Pred. No. 0.00057;
                                                                                                                                                                                                                                                                                                                                             100.0%; Preu. ...
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(UNAM ) UNIV VAN AMSTERDAM.
(UYAM-) UNIV AMSTERDAM ZIEKENHUI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 3; 7pp; English.
                                                                                                                        Disclosure; Page 3; 7pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                   1 RIORGPGRAFVTIG 14
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                                                                                                                                                                                                                                                                                                                                                                  14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          causing or related virus
              Meloen RH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Meloen RH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
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                                            WPI; 1989-108193/15
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                                                                                                                                                                                                                                                                                                     Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-OCT-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-AUG-2003
30-MAR-1992
              Goudamit J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAP95349;
                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                       field.)
                                                                                                                                                                                                                                                                                                                                                                    Matches
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The sequences given in AAR29237-43 represent a portion of the V3 loop region of the gpl20 envelope protein of various HIV isolates. These sequences can be used to define the specific isolate. All these viral varients exhibit complete homology at residues 7-11 of the given sequence and at least 36% homology with the remaining 12 amino acids of the sequence. Viruses containing these sequences are recognised by the heteroconjugate enzyme of the invention. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 24-OCT-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hetero-conjugate antibodies for treating HIV infections - comprise an antibody specific for an effector cell surface antigen and an antibody a V3 loop of GP-120 envelope protein of HIV.
                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
  S
blocked or modified otherwise. (Updated on 27-AUG-2003 to correct field.)
                                                                                                                                                                                                                                                                                                                                                                     V3 loop; gpl20; envelope protein; MN; prototype; virus; varient;
homology; heteroconjugate; enzyme; HIV.
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                                                                      Score 72; DB 1; Length 17;
Pred. No. 0.00057;
                                                                                                   0; Indels
                                                                                 100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus; IIIB varient
                                                                                                                                                                                                                                                                                                                                            V3 loop region epitope from IIIB isolate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 32; 69pp; English.
                                                                                                                                                                                                                               AAR29241 standard; peptide; 17 AA.
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                                                                       93.58;
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                                                                                                                              1 RIQRGPGRAFVTIG 14
                                                                                                                                                RIORGPGRAFVTIG 17
                                                                                                                                                                                                                                                                                                                (first entry)
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Matches 14; Conservative
                                                                                                   14; Conservative
                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Higgins PJ, Potts BJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (REPK ) REPLIGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-415475/50.
                                                                                       Similarity
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                                           Sequence 17 AA;
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                                                                                                                                                                                                                                                                                  24-OCT-2003
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                                                                       Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                         AAR29241;
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RESULT 125 AAR32407

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Human immunodeficiency virus 1
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                      WO9429339-A1.
                                                               08-JUN-1994;
                                                                                   199-JUN-1993;
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20-OCT-1997
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                                           22-DEC-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW25834;
                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 127
                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                       Nine peptides from the V3 regions of HIV-1 isolates IIIb, RF and MN were incorporated into terravalent multiple antigen peptide systems (MAPS) (see AAR32406-14). Parallel groups of three peptides with chain lengths spanning from 11-24 residues were synthesised in MAPS formate for each isolate. ELIS assays demonstrated that antisera titers in mice were closely related to the length of the IIIb peptide used for the immunisation - the longer the stronger the response. There was no substantial antibody prodn. in mice against the other two series of peptides, RF (B4-B6), and MN (B7-B9), except for a low reactivity in the gp; immunised with B8 (MN isolate). Specificity tests of the B cell response demonstrated that the T cell epitope (AAR2415) also serves as B cell epitope. (Updated on 25-MAR-2003 to correct PN field.)
                                                                    Sequence of peptide B2 which comprises AAs 312-281 from the V3 region of HIV-1 isolate IIIb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T-cell; epitope; HIV-1; core protein; p24E; B-cell; antigen; gp160; gag; pol; vaccine; multimeric peptide; AIDS; 3D organisation.
                                                                                                                                                                                                                                                                                                       New multiple antigen peptide(s) as HIV vaccines - include a dendritic core covalently bonded to peptide including the sequence IGPGR.
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93.5%; Score 72; DB 2; Length 17;
Best Local Similarity 100.0%; Pred. No. 0.00057;
Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T cell epitope derived from V3 isolate LAI
                                                                                                     HIV-1; vaccine; dendritic core; ss.
AAR32407 standard; peptide; 17 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR68664 standard; peptide; 17 AA.
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                                                                                                                                                                                        92WO-US006688.
                                                                                                                                                                                                            91US-00744281
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                          (revised)
(first entry)
                                                                                                                                                                                                                               (REPK ) REPLIGEN CORP. (UYRQ ) UNIV ROCKEFELLER.
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                                                                                                                                                                                                                                                                                   WPI; 1993-093730/11.
                                                                                                                                                                                                                                                              Tam JP, Profy AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 AA;
                                                                                                                                                                                      11-AUG-1992;
                                                                                                                                                                                                            13-AUG-1991;
                                                                                                                                              WO9303766-A1
                                                                                                                                                                    04-MAR-1993.
                                         25-MAR-2003
04-JUL-1993
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06-SEP-1995
                                                                                                                          Synthetic
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                     AAR32407;
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This sequence represents a T-cell epitope derived from the V3 sequence of the HIV-1 isolate LAI, which may be linked to a B-cell epitope from the V3 (MN) loop from HIV-1. These chimeric peptides may then be used in the production of HIV-1 vaccines. These peptides may then be used in the production of Multimeric peptides in which the peptides are C-terminally modified by the addition of a Lys residue which is modified on its epsilon amino acid to carry an additional copy of the peptide molecule. The linear and multimeric peptides may be used for the treatment of AIDS by acting to displace the binding of HIV virus to human carminal cells or by disturbing the 3D organisation of the virus. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 16-OCT-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                        Novel tandem synthetic HIV-1 peptide(s) - comprising T-cell epitope gag protein linked to B-cell epitope of V3 loop protein of an HIV-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93.5%; Score 72; DB 2; Length 17; 100.0%; Pred. No. 0.00057;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW25834 standard; peptide; 17 AA.
94WO-CA000317.
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                                                                   93US-00073378
                                                                                                                                                                                                     Sia CDY, Chong P, Klein MH;
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(first entry)
                                                                                                                                      (CONN-) CONNAUGHT LAB LTD
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Gaps

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us-08-869-386-1.rag

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New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 10; Page 119; 154pp; English.
                                             Disclosure; Col 21; 41pp; English
                                                                                                                                                                                                                                                                       AAW76848 standard; peptide; 17 AA
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                                                                                                                                                                                                  14; Conservative
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       WPI; 1997-332082/30.
                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus infection.
                                                                                                                                                                  Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                     Mus sp.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                            13-FEB-1998;
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immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gp120 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gp120 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as waccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gp120, so prevents or inhibits 'bystander' apoptosis of unifected T cells that are bound to gp120 protein, maximising induction of protective antiviral T cell immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |mmunogen; vaccine; HIV-1; T-cell; B-cell; epitope; core protein; gp120;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-cell epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitopes are generally designed based on the p24 core protein and the B-cell epitopes from the V3 loop of the gpl20 protein from various HIV-1 strains. This peptide represents the V3 loop epitope from the HIV-1 strain LAI. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                   93.5%; Score 72; DB 2; Length 17; 100.0%; Pred. No. 0.00057; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-1 strain LAI gp120 V3 loop epitope peptide.
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                                                                                                                                                                                                                                                                                                                                            Sequence 17 AA;
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                                                                                                                                                                                             The invention relates to new synthetic peptides comprising at least one amino acid sequence comprising an HIV gag protein T-cell epitope linked at its C or N-terminus to an amino acid sequence comprising a B-cell epitope of the V3 loop of an HIV env protein, which can be used to generate vaccines against HIV-1. The T-cell epitope sequence is pref. Palence from the T-helpper determinant core peptides P24E, P24N, P24L, P24M and P24H while the B-cell epitopes are derived from HIV strains including CTLB-56, V3MN, CTLB-29, CTLB-55, SF2, LAI, IIIB, RF, Z6, 2054, L14 and BX08. The peptides are chimaeric and can be linked to a branched Lys backbone. This sequence represents the B-cell env protein V3 loop peptide from HIV-1 strain LAI. (Updated on 25-MAR-2003 to correct PF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; waccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                     Tandem synthetic HIV peptide(8) useful as immunogens - comprising gag
protein T-cell epitope linked to env protein B-cell epitope.
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Pred. No. 0.00057;
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100.0%; Ex.
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Length 17;

93.5%; Score 72; DB 2; Le 100.0%; Pred. No. 0.00057;

Best Local Similarity

Query Match

Sequence 17 AA;

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WPI; 1999-550482/46
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                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 AA;
                                                                                                              06-JUN-1995;
                                                                                                                                09-JUN-1993;
                                                                          US5951986-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-JUN-2004
                                                                                                                                                                            Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADN14075;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADN14075
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         °;
                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a synthetic peptide comprising an amino acid sequence containing a T-cell epitope of an HIV gag protein linked a lts C terminus to an amino acid sequence containing a B-cell epitope of an HIV gp41 protein and containing the amino acid sequence: XILKDWAZ; where XI = E, A, G or Q, and X2 = A or T, or an amino acid sequence capable of eliciting an HIV-specific antiserum and recognizing the sequence XILKDWAZ. The synthetic peptide is useful in vaccines against HIV infection and in diagnostic applications. AAW98892 to AAW98996, and AAW98899 to AAW98999 represent synthetic peptides from the present
 Gaps
                                                                                                                                                                                                                                                                                                                                                    Synthetic chimeric HIV polypeptides - comprising gag protein T-cell epitope linked to gp41 B-cell epitope.
                                                                                                                                                       HIV-1; human immunodeficiency virus; vaccine; T-cell epitope; gag protein; B-cell epitope; gp41 protein; chimeric; infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 17;
0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 72; DB 2; Le
Pred. No. 0.00057;
0; Mismatches 0;
                                                                                                                                      HIV-1 vaccine synthetic peptide SEQ ID NO:35.
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                Example 1; Col 41-42; 41pp; English
                                                                                AAW99958 standard; peptide; 17 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93.5%; Sc.
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY39756 standard; peptide; 17 AA.
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                                                                                                                                                                                           Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                   93US-00073378.
                                                                                                                                                                                                                                                                                                                 Chong P, Klein MH, Sia CDY;
                                                                                                                                                                                                                                                  95US-00462507
                  1 RIORGPGRAFVTIG 14
                                  4 RIORGPGRAFVTIG 17
                                                                                                                                                                                                                                                                                               (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RIORGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 RIQRGPGRAFVTIG 17
                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                  WPI; 1999-189590/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
Les 14; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 AA;
                                                                                                                    05-MAY-1999
                                                                                                                                                                                                                                                  05-JUN-1995;
                                                                                                                                                                                                                                                                   09-JUN-1993;
                                                                                                                                                                                                              US5876731-A.
                                                                                                                                                                                                                                02-MAR-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-OCT-2003
26-NOV-1999
                                                                                                                                                                                   Synthetic
                                                                                                  AAW99958;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY39756;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Matches
                                                              RESULT 130
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY39756
ID AAY3
XX
AC AAY3
XX
DT 17-0
DT 26-N
                                                                        AAW99958
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a fragment of a HIVI protein, and can be used in the immunogenic composition of the invention. The composition comprises a synthetic fusion polypeptide which includes a sequence encoding 1 or more T cell epitopes and a sequence encoding 1 or more B cell epitopes and a sequence encoding 1 or more B cell epitopes and a proteins. The compositions are useful as vaccines against HIV infection. The composition induces HIV-1 specific polyclonal antibodies that are opsonising and antiviral. The peptide components may be selected to induce a response against different viral isolates and in subjects who recognise different T cell epitopes. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV; antigen; epitope; T cell; MHC; major histocompātibility complex; CTL; cytotoxic T lymphocyte; HIV infection; cancer; tuberculosis; tumour; hepatitis; melanoma; breast cancer; Hodgkin lymphoma; nasopharyngeal carcinoma; vaccine; immune response; hyaluronic acid; HA; CD8+ T cell; OD4+ T cell; viral infection; bacterial infection; fungal infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency virus, useful antigens in producing vaccines.
                                                                       HIV; vaccine; immunogenic composition; T cell epitope; B cell epitope;
infection; antibody; antiviral.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADN14075 standard; peptide; 17 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Col 22; 43pp; English.
                                                                                                                                                                                           Human immunodeficiency virus 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV helper T cell epitope #42.
HIV1 chimeric peptide V3-LAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chong P, Sia CDY;
                                                                                                                                                                                                                                                                                                                                                                                                                       95US-00467881.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93US-00073378.
94US-00257528.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 RIORGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 RIORGPGRAFVTIG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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Best Local Similarity 100.
Matches 14; Conservative
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The inversion relates to macutaring an immune system response to an antigen in a mammal comprising administering to the mammal a particle-free therapeutic comprising administering to the mammal a particle-free therapeutic comprising a hyaluronic acid (HA) polymer analogue covalently linked to at least one peptide that comprises a T cell epitope comprises a sequence of at least about eight amino acids of the antigen. Also included are a method of improving major histocompatibility complex (MHC) presentation of a T cell epitope of an antigen in a mammal (Comprising administering to the mammal the complex (MHC) presentation of a T cell epitope of an antigen in a mammal (Comprising administering to the mammal the complex (MHC) Class I molecule and by a CD8+ T cell of the mammal. Or an antibody that recognises the antigen. The immune system response comprises a cytotoxic T lymphocyte, a CD4+T CC collygate, where the antigen is an antigen of the conjugate, where the antigen is an antigen of a pathogenic agent or a tumour cell. The immune system response to the antigen is an antigen of a cumour cell. The immune system response to the antigen is an antigen of a gaministration of the conjugate, where the antigen is an antigen of a customer expense to an antigen in a mammal. The composition and methods are useful for modulating i.e. enhancing or diminishing, an immune coll importing major histocompatibility complex presentation of a T cell conjugates are useful as peptide vaccines against an antigen of a tumour cell) in a mammal. The polymeric hyaluronic acid conjugates are useful as peptide vaccines against an antigen, a conjugates are useful as peptide vaccines against an avoid is a conjugate and an antigen in a mammal. The polymeric conjugates are useful as peptide vaccine compositions are conjugated are useful as peptide vaccines against an avoid is a composition of the patential for treating or preventing diseases associated with any of the cantigens above e.g. HIV infection, cancer, tuberculosis, hepatitis administ
                                                                                                                                                                                                                                                                                                                                                                       Modulating an immune system response to an antigen in a mammal, comprises administering a particle-free therapeutic comprising a hyaluronic acid polymer analogue covalently linked to a peptide that comprises a T cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 additional adjuvants, but still induce a stronger cell-mediated response than peptide vaccines of the prior art. The present sequence is an HIV-1 derived epitope suitable for the vaccine of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to modulating an immune system response to an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 12; 23pp; English.
                                                    05-FEB-2002; 2002US-00062710.
                                                                                                      08-AUG-2001; 2001US-0310498P
                                                                                                                                                                                                                                                                   Chu Y, Qiu J;
                                                                                                                                                                                                                                                                                                                  WPI; 2003-540464/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 AA;
                                                                                                                                                          (LIFQ/) LI P Q.
                                                                                                                                                                                   (CHUY/) CHU Y.
(QIUJ/) QIU J.
13-MAR-2003.
                                                                                                                                                                                                                                                                 Li FO,
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Gaps ö Length 17; 0; Indels 93.5%; Score 72; DB 7; Le 100.0%; Pred. No. 0.00057; tive 0; Mismatches 0; 2 IQRGPGRAFVTIGK 15 1 IQRGPGRAFVTIGK 14 14; Conservative Local Similarity Query Match **fatches** 셤 ઠે

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ADR18895 standard; peptide; 17 AA. ADR18895; RESULT 133 ADR18895 8XXX

04-NOV-2004 (first entry)

three-dimensional atomic structural conformation; HIV-1 V3 loop related peptide SEQ ID NO:28.

protein co-ordinate data; V3 loop peptide; HIV-1; envelope glycoprotein; gpl20; human monoclonal antibody 447-52D; nurine monoclonal antibody 0.5 beta; immunogen; immunogenic; loop epitope; HIV-1 infectivity inhibitor; anti-HIV; vaccine; HIV-1 infection 

Human immunodeficiency virus 1.

Synthetic.

WO2004069863-A2

19-AUG-2004.

34-FEB-2004; 2004WO-US003304.

04-FEB-2003; 2003US-0444682P.

(UYNY ) UNIV NEW YORK STATE. (YEDA ) YEDA RES & DEV CO LTD.

Anglister J, Sharon M, Schapira M, Zolla-Pazner S, Rosen O;

NPI; 2004-625447/60.

Composition for inhibiting HIV-1 infection, comprises isolated peptide molecule that mimics atomic structural conformation of V3 loop peptide of HIV-1 envelope glycoprotein that is bound to, and constrained by human monoclonal antibody.

Claim 8; SEQ ID NO 28; 127pp; English.

The present invention describes a composition (C1) which comprises an isolated peptide molecule or isostere that minics the three-dimensional isolated peptide molecule or isostere that minics the three-dimensional conformation of the V3 loop peptide of the HIV-1 and constrained by, human converge glycoprotein gp120 that is bound to, and constrained by, human conformal antibody (MAD) 447-520, marine MAD 0.5 beta or an antigen binding fragment of the MAD, where the constrained V3 loop peptide different of conformation from the same V3 loop peptide when it is in free form. Also described: (1) identifying (MI) from several existing compounds a molecule that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (3) a composition (C2) that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 antibody response specific for a V3 loop immunogen or as an inhibitor of that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of or induction of an anti-HIV-1 antibody response specific for a V3 loop for induction of an anti-HIV-1 antibody response specific for a V3 loop for induction of an anti-HIV-1 antibody response specific for a V3 loop or X4 co-receptor and thereby inhibiting HIV-1 infectivity. Comprising cc mobal representing the conformationally constrained by NG; and a 3D made of a constrained HIV V3 view peptide; (7) a computer readable medium (CM) computing platform for generating composition of a valope peptide that is bound to 447-520 or 0.5beta MAD or its antigen composition and carrier or excipient; (6) a computer readable medium (CM) comprising fragments, computing platform for a valop peptide that is conformationally constrained by being bound to 447-520 or 0.5beta MAD or its antigen beinding of HIV-1 to chemokine receptor/HIV-1 co-receptor. (C1) is useful for invivo inhibition of HIV-1 infection

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Gaps

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us-08-869-386-1.rag

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93.5%; Score 72; DB 2; Length 18; 100.0%; Pred. No. 0.0006; ive 0; Mismatches 0; Indels
                                                                                                              AAW03404 standard; peptide; 18
                                                            RIORGPGRAFVTIG 18
                                        1 RIORGPGRAFVTIG 14
                      14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1994-128412/16.
           Similarity
                                                                                                                                                                                                                                                          Key
Modified-site
                                                                                                                                                                                                                                                                                                                                         Modified-site
                                                                                                                                                                                                                                                                                                                                                           Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                            12-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-OCT-1992;
                                                                                                                                                       10-0CT-1996
                                                                                                                                                                                                                                                                                                                                                                                                      GB2271995-A.
                                                                                                                                                                                                                                                                                                                                                                                                                         14-MAY-1994
                                                                                                                                                                                                                                       Synthetic
                                                                                                                                  AAW03404;
  Query Match
           Best Local
Matches 1
                                                                                                      AAW03404
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in an infected subject. The present sequence represents a peptide used in the exemplification of the present invention.
                                                                                                                                                                                                                                                       Cyclic, HIV; principle neutralising determinant; PND; conjugate; immunological carrier; antipeptide; anti-HIV; HIV-neutralising; antibody; vaccine; immune response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequences given in AAR38520-26 are cyclic HIV principle neutralising determinant (PND) peptides. They are stable compounds which, when conjugated with an immunological carrier, may be used to raise antipeptide, anti-HIV and HIV-neutralising antibodies. These may be used in vaccines for prevention or treatment of HIV-related diseases. They may also be used as reagents in the study of structure/function relationships in induction of HIV-neutralising immune response. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                    /note= "C-terminal condensed to the N-terminal of Lys1 to form cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cyclic HIV principal neutralising determinant (PND) peptide(s) - useful as vaccines for treating and preventing HIV-mediated diseases e.g. AIDS
                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                        /note= "Alpha amino group forms a cyclic peptide by condensing to the C-terminal of Gly18"
                                                                       ò
                                                   Length 17;
                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                 note= "Abu = gamma aminobutyric acid"
                                                  DB 8; Le
                                           93.5%; Scorio 100.0%; Pred. No. 0, Mismatches
                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 26; 27pp; English.
                                                                                                                                                              AAR38526 standard; peptide; 18
                                                                                                                                                                                                                                    Cyclic HIV PND peptide cPND535
                                                                                                                                                                                                                                                                                                                                                                      /label= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92EP-00300243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91US-00807943
                                                                                         1 RIORGPGRAFVTIG 14
                                                                                                      4 RIQRGPGRAFVTIG 17
                                                                                                                                                                                                                 (first entry)
                                                                      14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tolman RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1993-228557/29.
                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                              Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                     Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                          Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-DEC-1991;
                                                                                                                                                                                                       25-MAR-2003
21-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                  EP551689-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUL-1993
                                                                                                                                                                                   AAR38526;
                                                                                                                                                                                                                                                                                                 Synthetic
                                                  Query Match
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                                                                                                                                           RESULT 134
                                                             Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New conjugates of outer membrane protein and HIV epitope - for generating HIV-neutralising response, have components joined by anionic spacer to ensure solubility of prod.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A new conjugate immunogen comprises (a) the OMPC of Neisseria meningitidis b as a protein carrier, (b) a principal neutralizing determinant (PND) of HIV as a peptidyl epitope against which immune responses are desired, and (c) a low mol. wt. anionic spacer linking (and (b). The conjugate is water-soluble, yet can carry a high peptide epitope loading. It is useful as a vaccine against HIV. The present sequence is an example of a PND used in the conjugate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "the carboxy group is condensed onto an amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                  conjugate; PND; HIV; principal neutralizing determinant; OMPC; outer membrane protein complex; anionic spacer; vaccine; human immunodeficiency virus; water-soluble.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lombardo VK;
                                                                     HIV principal neutralizing determinant cPND535
                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Leanza WL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lys(1). See above"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 10; Page 70; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                   label= cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93GB-00020943.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   label= Abu
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MERI ) MERCK & CO INC
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Sequence 18 AA;

us-08-869-386-1.rag

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The present invention describes a composition (C1) which comprises an isolated peptide molecule or isostere that mimics the three-dimensional conformation of the V3 loop peptide of the HIV-1 envelope glycoprotein gpl20 that is bound to, and constrained by, human monoclonal antibody (MAb) 447-520, murine MAb 0.5 beta or an antigen binding fragment of the MAb, where the constrained V3 loop peptide of differs in conformation from the same V3 loop peptide when it is in free form. Also described: (1) identifying (M1) from several existing compounds a molecule that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (2) designing a molecule that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (3) a composition (C2) inhibitor of binding of HIV-1 v3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor on the surface of a receptor-bearing target cell; (3) a composition (C3) contraining (C1) and an excipient; (5) a pharmaceutical composition (C4) useful for blocking the interaction of HIV-1 with an R5 corpustion of an anti-HIV-1 antibody response specific for a V3 loop epitope, comprising (C1) and an excipient; (6) a computing platform for generating (C1) and a carrier or excipient; (6) a computing platform for generating corpus enting the conformationally constrained structure of a V3 corpuse model representing the conformationally constrained structure of a V3 corpuse for bound to 447-520 or 05beta MAb or its antigen binding fragments, compusiting a 3D atomic structure defined by NC; and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Composition for inhibiting HIV-1 infection, comprises isolated peptide molecule that mimics atomic structural conformation of V3 loop peptide of HIV-1 envelope glycoprotein that is bound to, and constrained by human monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              three-dimensional atomic structural conformation; protein co-ordinate data; V3 loop peptide; HIV-1; envelope glycoprotein; gp120; human monoclonal antibody 447-520; murine monoclonal antibody 0.5 beta; immunogen; immunogenic; V3 loop epitope; HIV-1 infectivity inhibitor; anti-HIV; vaccine; HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anglister J, Sharon M, Schapira M, Zolla-Pazner S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV-1 V3-IIIB peptide SEQ ID NO:4.
                                                                                                                                                                                                                                                                        ADR18878 standard; peptide; 18 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYNY ) UNIV NEW YORK STATE.
(YEDA ) YEDA RES & DEV CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-FEB-2004; 2004WO-US003304.
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1 RIQRGPGRAFVTIG 14
                                            04-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-625447/60.
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Rosen O;

data that includes a set of structure coordinates defining a 3D structure of a V3 loop peptide that is conformationally constrained by being bound to 447-52D or 0.5beta MAD or its antigen binding fragment. (C1) has antition 447-52D or 0.5beta MAD or its antigen binding fragment. (C1) has antitivities, and can be used in vaccines, and as an inhibitor of binding of HIV-1 to chemokine receptor/HIV-1 co-receptor. (C1) is useful for in vivo inhibition of HIV-1 infection. (C1) or (C2) is useful for producing a medicament utilised for treating or preventing HIV-1 infection. (C3) or (C4) is useful for inducing in a subject an anti-HIV-1 neutralising antibody response specific for a V3 loop epitope. (C4) is useful for preventing an HIV-1 infection in an uninfected subject at risk for such infection or for inhibiting viral spread and disease progression in an infected subject. The present sequence represents a peptide used in the exemplification of the present invention. The peptide sequence given represents the sequence from amino acid 310-328 of the external protein gp120 from HIV. This comprises a disulphide loop in the third variable region. It was used in an example of the invention and was incorporated into hepatitis B surface antigen (HBsAg) particles. The hybrid formed in this reaction is useful as a vaccine for the prophylactic treament of various infectious diseases eg. AIDS. Conjugation of this peptide with the HBsAg particle allows its processing to be directed via a non-endosomal route. In this way the gp120 fragment can be recognized by cytotoxic lymphocytes. (Updated on 25-MAR-2003 to correct PN field.) New immunogenic hybrid polypeptide(s) for vaccine formulations - compris S antigen of hepatitis B linked via spacer to heterologous antigen, e.g. gD from HSV or gpl20 form HIV. Gaps ö Hepatitis B; surface antigen; AIDS; cytotoxic lymphocytes; disulphide loop; variable region. 93.5%; Score 72; DB 8; Length 18; 100.0%; Pred. No. 0.0006; tive 0; Mismatches 0; Indels Van Wijnendale F, Baijot M, Prieels J; (SMIK ) SMITHKLINE BEECHAM BIOLOGICALS. Disclosure, Page 28; 38pp; English. AAR25471 standard; protein; 20 AA. 91WO-EP002422 90GB-00027623 91GB-00005993. 1 RIORGPGRAFVTIG 14 s kiokgpckarvric 18 (revised)
(first entry) Query Match Best Local Similarity 100. Matches 14; Conservative WPI; 1992-250032/30. 73 loop structure. Sequence 18 AA; WO9211291-A1 16-DEC-1991; 20-DEC-1990; 21-MAR-1991; 25-MAR-2003 15-JAN-1993 09-JUL-1992. Synthetic. AAR25471; RESULT 137 AAR25471 ઠે 셤 

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Claim 10; Page 119; 154pp; English.
                                                                 AAW76842 standard; peptide; 20 AA.
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   Query Match
Best Local Similarity 100.00
Conservative
Conservative
                              1 RIORGPGRAFVTIG 14
                                       7 RIORGPGRAFVTIG 20
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Best Local Similarity 100.0
Matches 14; Conservative
                                                                                                                                                                                                            Zambidis E;
                                                                                                                                                                                                                      WPI; 1998-506315/43.
Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                Sequence 20 AA;
                                                                                                                                          sp.
Sapiens.
                                                                                      25-JAN-1999
                                                                                                                                                                             13-FEB-1998;
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                                                                                                                                                        WO9836087-A1
                                                                                                                                                                  20-AUG-1998.
                                                                            AAW76842;
                                                                                                                                                                                                            Scott D,
                                                        RESULT 138
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The present invention describes an anti-human leukocyte antigen (HLA) antibody (I) useful for the preparation of a medicament for treating a disease involving a proliferative immune response. (I) has anti-HIV, antiinflammatory, neuroprotective and cytostatic activities, and can be used in vaccines and in gene therapy. The antibody (I) is useful for the preparation of a medicament for treating diseases involving a proliferative immune response, e.g. HIV, inflammatory diseases, autoimmune diseases, axonal or nerve damage or related impairment or cancers, and other diseases or conditions with an inflammatory component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents an HIV gpl20 V3 loop derived peptide, which is used in the exemplification of the present invention. (Updated on 23-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                     Anti-human leukocyte antigen antibody; anti-HLA antibody; anti-HIV; proliferative immune response; antiinflammatory; neuroprotective; cytostatic; gene therapy; vaccine; inflammatory disease; nerve damage; autoimmune disease; axonal damage; cancer; inflammatory; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Use of anti-HLA antibody for the preparation of a medicament for tradisease involving a proliferative immune response e.g. HIV, inflammatory diseases, autoimmune diseases, axonal/nerve damage or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dalgleish AG, Cadogan M, Heeney J, White SDT;
                                                                                                                                                                                                                                                                                                                                                     HIV gp120 V3 loop derived peptide ARP740/28.
                                                                                                                                                                       ABP57070 standard; peptide; 20 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 41; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    32-JUL-2002; 2002WO-GB003037.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JUL-2001; 2001GB-00016151.
29-NOV-2001; 2001GB-00028638.
28-JAN-2002; 2002GB-00001896.
28-MAR-2002; 2002GB-00007509.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               related impairment, cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93.5%;
  14
                          7 RIORGPGRAFVTIG 20
                                                                                                                                                                                                                                                                              (revised)
(first entry)
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1 RIORGPGRAFVTIG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 AA;
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14-APR-2003
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                                                                                                                                                                                                                            ABP57070;
                                                                                                                         RESULT 139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, so prevents or inhibits 'bystander' apoptosis of unifected T cells that are bound to gpl20 protein, maximising induction of protective antiviral T cell immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, 1gH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and anintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                       Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #12.
                                                            93.5%; Score 72; DB 2; Length 20; 100.0%; Pred. No. 0.00066; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98WO-US002766.
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             Length 20;
            93.5%; Score 72; DB 6; Le 100.0%; Pred. No. 0.00066; ive 0; Mismatches 0;
                                                          1 RIORGPGRAFVTIG 14
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Query Match
Best Local Similarity 100...
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AAR66425
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Gaps

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93.5%; Score 72; DB 2; Length 20; 100.0%; Pred. No. 0.00066; tive 0; Mismatches 0; Indels

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(USSH ) US SEC DEPT HEALTH.
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                                                                                                                                                             WPI; 1995-006707/01.
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 AA;
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                               WO9426785-A1
                                                                          13-MAY-1994;
                                                                                              14-MAY-1993;
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03-AUG-1995
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                                                    24-NOV-1994.
           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR66429;
                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a val at position 11 and substms. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substms. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-10, the Phe residue at position 10 in peptide 18 has been replaced by a 11e residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                   Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                             T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                                                                                                                           Pendleton CD, Nara P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92.2%; Score 71; DB 2; I 93.3%; Pred. No. 0.00072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 33; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR66420 standard; peptide; 15 AA.
 AAR66425 standard; peptide; 15 AA
                                                                                                                                                                                                          94WO-US005142.
                                                                                                                                                                                                                               9303-00060988
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                                          (revised)
(first entry)
                                                                                                                                                                                                                                                                           Berzofsky JA, Ahlers JD,
                                                                         HIV-1 IIIB peptide 18-10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 93.3
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                                                                                                                                                                                                                                                                                               WPI; 1995-006707/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 AA;
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                                          25-MAR-2003
03-AUG-1995
                                                                                                                                          Synthetic.
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                     AAR66425;
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AAR66420
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Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising series from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-AR6430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-5, the Gly residue at position 5 in peptide 18 has been replaced by an Ala residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                         Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                                                                                        Shirai M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92.2%; Score 71; DB 2; Length 15; 93.3%; Pred. No. 0.00072; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                        Berzofsky JA, Ahlers JD, Pendleton CD, Nara P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 33; 120pp; English
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94WO-US005142.
                                                                              93US-00060988.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIORAPGRAFVTIGK 15
                                                                                                                                                             (USSH ) US SEC DEPT HEALTH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 93.3
Matches 14; Conservative
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Nara P, Shirai M;

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Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66410) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a val at position 11 and substnes at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of determinant sequence (FGRAF). In peptide 18-14, the Gly residue at position 11 in peptide 18 has been replaced by an Ala residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                   Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   92.2%; Score 71; DB 2; Length 15; 93.3%; Pred. No. 0.00072; ive 0; Mismatches 1; Indels
                                   Pendleton CD,
                                                                                                                                                                                                                                                                                  Example 1; Page 33; 120pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ahlers JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14; Conservative
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                                                                                              WPI; 1995-006707/01.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 AA;
                                   Berzofsky JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9426785-A1.
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03-AUG-1995
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AAR66423
XX
AC AAR6641
XX
AC AAR6641
DT 25-MAR.
DT 25-MAR.
DT 25-MAR.
DT 25-MAR.
CO 33-AUG.
CO 31-AUG.
CO 31-A
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Gaps ö

AAR66423 standard; peptide; 15 AA.

T cell helper site, cytotoxic T cell response, neutralising antibody, human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.

(USSH ) US SEC DEPT HEALTH

Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV. WPI; 1995-006707/01.

Shirai M;

Nara P,

Ahlers JD, Pendleton CD,

Berzofsky JA,

Example 1; Page 33; 120pp; English.

Single residues from the HIV-1 RF sequence (AAR66415) replaced residues

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in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals fimmunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-R66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRARP). In peptide 18-8, the Arg residue at position 8 in peptide 18 has been replaced by an Ala residue. (Updated on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-7, the Gly residue at position 7 in peptide 18 has been replaced by an Ala residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                                                                                                                        92.2%; Score 71; DB 2; Length 15; 93.3%; Pred. No. 0.00072; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nara P,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR66422 standard; peptide; 15 AA.
                                                                                                                                                                                                 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                             (USSH ) US SEC DEPT HEALTH.
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(first entry)
                                                                                                                                                                                                                                                                                                                14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1 IIIB peptide 18-7.
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                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                    Sequence 15 AA;
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03-AUG-1995
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AAR66421;

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The invention relates to a novel monoclonal antibody (MAb) NM03 which binds to epitopes from the human immunodeficiency virus type 1 (HIV-1). The antibody was raised conventionally by immunising Balb/c mice with purified live HIV-1 MN, then isolating splencytes and fusing them to P3- K63-Ag8-U1 cells. Hybridomas were then screened with membrane fractions from infected and non-infected H9 cells. The MAb was observed to bind to a protein band of 120 kD on a Western blot of separated, denatured HIV-1 proteins. This binding was shown to be between residues 320-327 by proteins. This binding was shown to be between residues 320-327 by contope mapping by ELISA and competitive binding. The ability of the MAb to invention of cells by HIV-1 shown by infecting H9 cells with live strains of HIV-1 and testing infection by a p24 assay. This peptide sequence represents the V3 loop region from HIV-1 strain IIIB, where the NAD NOAD binds. The MAb can be used for the passive immunisation of thumans susceptible to, or infected with HIV-1. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                Epitope, human immunodeficiency virus type 1, HIV-1; gp120; gp160; monoclonal antibody; V3 loop; immunisation; mouse; splenocyte; hybridoma; membrane fraction; passive immunisation; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monoclonal antibody specific for human immunodeficiency virus type 1 MN strain - for passive immunisation against infection.
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100.0%; Pred. No. 0.0018;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                     HIV-1 strain IIIB gp120 V3 loop peptide.
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                                         AAW22327 standard; peptide; 13 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1
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hes 13; Conservative
                                                                                                                                                                        (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUL-1994;
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                                                                                                                                                                     17-OCT-2003
18-SEP-1997
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                                                                                                       AAW22327;
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AAW 2337

AAW 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
                                                                                                                                         Gaps
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                                                                    92.2%; Score 71; DB 2; Length 15; 93.3%; Pred. No. 0.00072; ive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR66421 standard; peptide; 15 AA.
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                                                                                                                                         14; Conservative
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                                                                 Query Match
Best Local Similarity
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      Sequence 15 AA;
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03-AUG-1995
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Gaps

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Monoclonal antibody; HIV-1gp120; gp160; infection; H9 cell;

Query Match

Best Loca Matches

RESULT 146

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Polypeptide inducing helper T cell, cytotoxic T cell and antibodies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-MAY-1993;
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03-AUG-1995
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                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 149
                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Glutamic acid; glutamine; vaccine; major histocompatibility complex; MHC; immunomodulator; antiallergic; endocrine; neuroprotectant; virucidal; pactericidal; antiparasitic; fungicidal; cytostatic; medicine; pharmaceutical; immune dozorder; immune disorder; autoimmune; hypersensitivity; allergy; graft rejection; infection; hormonal disorder; central nervous system disease; cancer; melanoma; anti-melanoma vaccine;
                                                                                                                                                       Monoclonal antibody which binds to HIV-1gp120 or gp160 - used to prevent and treat human HIV infection.
                                                                                                                                                                                                    AAW62889-900 represent peptides used to identify a peptide sequence (AAW62874) present in HTV-1gp120 or gp160 which is bound by the monoclonal antibody of the invention. The antibody neutralises in vitro the infection of H9 cell by an active HIV strain MN according to the p24 analytical method. The antibody is used for treatment of human HIV
                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                          88.3%; Score 68; DB 2; Length 13; 100.0%; Pred. No. 0.0018; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vaccine related MHC ligand peptide SEQ ID NO:536.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Goetsch
HIV strain MN; treatment; human HIV infection.
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                                                                                                                 (NISP ) NISSIN SHOKUHIN KAISHA LTD.
                                                                                                                                                                                   Example 3; Page 8; 12pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                 AAM99433 standard; peptide; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1.
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                                                                            96JP-00344904
                                                                                                96JP-00344904
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(first entry)
                                                                                                                                                                                                                                                                                                                                  3 QRGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                    Local Similarity 100.
nes 13; Conservative
                                                                                                                                    WPI; 1998-433774/37
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                                                                                                                                                                                                                                                                         Sequence 13 AA;
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                                      JP10182489-A
                                                                            25-DEC-1996;
                                                                                               25-DEC-1996;
                                                         07-JUL-1998
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07-DEC-2001
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                   Synthetic
                                                                                                                                                                                                                                                       infection
                                                                                                                                                                                                                                                                                                                                                                                                                     AAM99433;
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The present invention describes a pharmaceutical compound (I) that contains an N-terminal glutamic acid (Glu) or glutamine (Gln) residue in the form of an addition salt with a strong, physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition on the form of an addition alt with a strong, physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition of (I) containing at least one (I) a vaccine containing at least one (I) containing at least one (I) and (a) a pharmaceutical with the presence of (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process of for preparing (I). (I) has immunomodulator, endocrine, antiallargic, containing acidil and cytostatic activities (I) are useful, in human or veterinary medicine, contained activities (I) are useful, in human or veterinary medicine, in pharmaceutical compositions (for treating immune disorders, e.g. impunction, infection, hormonal disorders and central nervous system diseases), also, where (I) is a MHC ligand (Ia), in vaccines for creatment or prevention of: (I) viral, bacterial, parasitic or fungal infections; or (ii) of cancers A particular application is in anti-melanoma vaccines. (I) are also useful for in vitro diagnosis of diseases of massociated with interactions between MHC and (I), e.g. melanoma and human immunodeficiency virus infection. AMM9888 to AAM9592 represent invention. We will impunct in pharmaceutical compounds from the present invention. CC (Updated on II-SEP-2003 to standardise OS field)
Stabilized pharmaceutical containing N-terminal glutamic acid or glutamine, useful e.g. in anti-melanoma vaccines, is an addition salt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pendleton CD,
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                                                                                                                                                                       Claim 9; Page 122; 149pp; French.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Berzofsky JA, Ahlers JD,
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                                                 glutamine, useful
with strong acid.
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single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue to the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR6640) and AAR6641). Analysis of the subscituted peptides (AAR66416-tyrosine was substd. for a Val at position 11 and substruct when a tyrosine was reduced when substns. were made in the principal neutralising sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-2, the Ile residue at position 2 in peptide 18 has been deleted. (Updated on 25-MAR-2003 to correct PN field.)
- to target antigen in hosts of different MHC haplotypes, esp
                       for therapeutic or prophylactic vaccines against HIV.
                                                                         Example 1, Page 33; 120pp; English.
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Sequence 14 AA;

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Gaps
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88.3%; Score 68; DB 2; Length 14; 100.0%; Pred. No. 0.0019; ive 0; Mismatches 0; Indels
                                  13; Conservative
                   Best Local Similarity
   Query Match
                                    Matches
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3 ORGPGRAFVTIGK 15 2 ORGPGRAFVTIGK 14 名 ઠ

AAW76897 standard; peptide; 15 AA.

AAW76897;

25-JAN-1999 (first entry)

Fusion immunoglobulin heavy chain HIV gp120 T cell epitope #16.

B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxie; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity. AAW76897

ID AAW7

XX AC AAW7

XX AC AAW7

XX B CS

YX B CS

XX CS

YX CS

Ношо варіелв gb. Mue

WO9836087-A1

20-AUG-1998

13-FEB-1998;

97US-0040581P. 13-FEB-1997;

98WO-US002766.

(AMNA-) AMERICAN NAT RED CROSS

Zambidis E; Scott D,

WPI; 1998-506315/43.

New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency virus infection

Claim 11; Page 120; 154pp; English.

The present invention describes a pharmaceutical compound (1) that contains an N-terminal glutamic acid (Glu) or glutamine (Gln) residue in the form of an addition salt with a strong, physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition containing at least one (I); (b) a vaccine containing at least one (I); where this is a major histocompatibility complex (WHC) ligand (Ia); (c) a method for in vitro diagnosis of diseases associated with the presence of (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process for preparing (I). (I) has immunomodulator, endocrine, antiallergic, neuroprotectant, virucidal, bactericidal, antiparasitic, fungicidal and

Claim 9; Page 63; 149pp; French

This sequence is an epitope used in the construction of a novel fusion immunosolobulin heavy chain (1g91) protein with a mammalian, especially human, 1gH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gp120 epitopes. Fusion Ig proteins and/or

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3lutamic acid; glutamine; vaccine; major histocompatibility complex; MHC;
transfected cells are used to tolerise subjects to gp120 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as accines, chemcherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gp120, so prevents or inhibits 'bystander' apoptosis of unifected T cells that are bound to gp120 protein, maximising induction of protective antiviral T cell immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunomodulator; antiallergic; endocrine; neuroprotectant; virucidal;
bactericidal; antiparasitic; fungicidal; cytostatic; medicine;
bacmaceutical; immune disorder; immune deficiency; autoimmune;
hypersensitivity; allergy; graft rejection; infection; hormonal disorder;
central nervous system disease; cancer; melanoma; anti-melanoma vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stabilized pharmaceutical containing N-terminal glutamic acid or glutamine, useful e.g. in anti-melanoma vaccines, is an addition salt
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                                                                                                                                                                                                                  88.3%; Score 68; DB 2; Length 15;
100.0%; Pred. No. 0.0021;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vaccine related MHC ligand peptide SEQ ID NO:186.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Klinguer-Hamour C, Corvaia N, Beck A,
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                                                                                                                                                                                                                                                                                                                                                                                                     AAM99083 standard; peptide; 15 AA.
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                                                                                                                                                                                                                      88.3%;
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                                                                                                                                                                                                                                     Local Similarity 100.
nes 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             with strong acid.
                                                                                                                                                                                      Sequence 15 AA;
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Matches
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in pharmaceutical compositions (for treating immune disorders, e.g. immune deficiency, autoimmune states, hypersensitivity, allergy, graft rejection, infection, hormonal disorders and central nervous system diseases), also, where (I) is a MHC ligand (Ia), in vaccines for treatment or prevention of: (i) viral, bacterial, parasitic or fungal infections; or (ii) of cancers. A particular application is in antimelanoma vaccines. (I) are also useful for in vitro diagnosis of diseases associated with interactions between MHC and (I), e.g. melanoma and human immunodeficiency virus infection. AAM988989 to AAM99592 represent peptides which can be used in pharmaceutical compounds from the present invention. (Updated on 11-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic peptides AAM63049-85 are used, with superantigens, to exemplify the invention. The specification describes a method for treatment of cancer which comprises incubating lymphocytes with a tumour-specific antigen or autoimmune disease related antigen and a superantigen. The treated cells are then introduced into the patient. The superantigen and the tumour-specific antigen or autoimmune disease related antigen and conjugated together. The products are used to treat cancer (carcinoma melanoma, lymphoma etc.), infections (bacterial, viral or eukaryotic) and autoimmune disease (e.g. idiopathic thrombocytopaenic purpura, rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antigen -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conjugates and polymers containing superantigen and therapeutic antigen for treatment of cancer, infection, autoimmune disease and graft rejection, also treatment by administering lymphocytes treated in vitro
  cytostatic activities. (I) are useful, in human or veterinary medicine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Superantigen, treatment, cancer, tumour-specific antigen, autoimmune disease related antigen, infection; bacterial, viral, eukaryotic, autoimmune disease; inhibit; pathological response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus type 1 (HIV 1) Env peptide 312-327.
                                                                                                                                                                                                                                                                                                                                      ;
                                                                                                                                                                                                                                                                                                   Length 15;
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100.0%; Pred. No. 0.0021;
iive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW63062 standard; peptide; 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-0044074P
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                                                                                                                                                                                                                                                                                                                                                                           3 ORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                2 ORGPGRAFVTIGK 14
                                                                                                                                                                                                                                                                                                                                    13; Conservative
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                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        by these antigens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immune response.
                                                                                                                                                                                                                                                        Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9826747-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Terman DS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW63062;
                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IGH) protein with a mammalian, especially human, IGH chain fused in frame at its N-terminus to one or more human immuno deficiency virus (HIV) gpl20 epitopes. Fusion IG proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autcommune disease. Induction of tolerance suppresses production of antibodies against gpl20, are bound to gpl20 protein, maximising induction of protective antiviral T cell immunity
arthritis, systemic lupus erythematosus, multiple sclerosis etc.). The antigens either induce an immune response or inhibit a patholgical response
                                                                                                                                                                                                                                                                                                                                                                                                                                           B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; yaccine; chemotherapy; immune response; modifier; tumour; microbial infection, autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                         Fusion immunoglobulin heavy chain HIV gp120 T cell epitope #58.
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                                                                                                           Length 18;
                                                                                                                                          0; Indels
                                                                                                       88.3%; Score 68; DB 2; 1
100.0%; Pred. No. 0.0024;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                    AAW79180 standard; peptide; 21 AA.
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                                                                                                                                                                             3 ORGPGRAFVTIGK 15
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                                                                                                                          Similarity 100.
13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scott D, Zambidis E;
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                                                                      Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                        25-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus sp.
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-FEB-1998;
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Length 21;

88.3%; Score 68; DB 2;

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20 so prevents or inhibits 'bystender' apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral
                                                                                                                                                                                                                            B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; igH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                      Gaps
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                                                                                                                                                                                                     Fusion immunoglobulin heavy chain HIV gp120 T cell epitope #20.
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Pred. No. 0.0028;
0; Mismatches 0; Indels
                      Indele
Pred. No. 0.0028;
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                                                                                                                               AAW76901 standard; peptide; 21 AA
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100.0%; Pre-
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        100.08;
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                                             3 ORGPGRAFVTIGK 15
                                                                    1 ORGPGRAFVTIGK 13
                     13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zambidis E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-506315/43.
        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       virus infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                   13-FEB-1997;
                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Scott D,
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The peptide represented in this sequence, and those represented by sequences AAR74604-7 are epitopes of HIV-1 gp120 that are effective to generate antibodias that inhibit infection of mucosal cells by HIV-1. These peptides are administered to the epithelium in a vaccine, or are used to generate mucosal antibodies and thereby inhibit infection by HIV-1. These peptides are used for inhibiting the entry of HIV into vaginal and rectal mucosal epithelium. The antibodies that can be generated from them are able to block subsequent infection by HIV. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1 gp120 peptide to inhibit mucosal epithelium cell infection – useful
in peptide vaccine to inhibit HIV-1 infection of vaginal or rectal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV isolates HIV-IIIB and HIV-RF; hybrid peptide PR137; therapy; AIDS; principal neutralising domain; antibodies; diagnosis; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gapa
                                                                                                                                                                                                                    HIV-1; HIV; AIDS; gp120; mucosal cell; epithelium; vagina; rectum; antibody; mucosal administration; vaccine; infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Svennerholm B, Vahlne A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88.3%; Score 68; DB 2; Length 24; 93.3%; Pred. No. 0.0031; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus hybrid peptide RP137.
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AAR74608 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR04475 standard; protein; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 2; Page 23; 34pp; English.
                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   94WO-US012152.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93US-00143577.
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(revised)
(first entry)
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                                                                                                                                                                           HIV-1 gp120 peptide #5.
                                                                                                  (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SYNT-) SYNTELLO INC.
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25-MAR-2003
20-SEP-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                   W09511701-A1
                                                                                                  16-OCT-2003
04-JAN-1996
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                                                 AAR74608;
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Variable region V3, found in the envelope protein gp120 of an AIDS or ARC causing or related virus.
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                                                                                                                                                                                 Synthetic human immunodeficiency virus vaccine - comprising hydrophilic peptide corresponding to at least 1 antigenic determinant of envelope glyco:protein recognised by B lymphocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligopeptide(s) corresp. to beta-turn variable region of gp.120 - used for diagnosis of and prodn of vaccines against AIDS and ARC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87.0%; Score 67; DB 2; Length 15; 93.3%; Pred. No. 0.0029; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                       Disclosure, Page 27; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vaccine; AIDS; ARC; HIV; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAP95356 standard; peptide; 17 AA.
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(UYAM-) UNIV AMSTERDAM ZIEKENHUI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 3; 7pp; English.
                      95US-00546515.
96US-00599266.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RIORGPGRAFVTIGK 15
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(first entry)
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                                                                                                             Haynes BF, Palker TJ;
                                                                                                                                                 WPI; 1997-244862/22.
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                                                                         (UYDU-) UNIV DUKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 AA;
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                      20-OCT-1995;
09-FEB-1996;
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30-MAR-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 셤
                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide RP137 comprises segments of the Principal Neutralising Domain (envelope protein) from isolates HIV-RF and HIV-IIIB. The last Cys carsidue is added for the purpose of crosslinking to carrier proteins. Cysteine residues may be added, so that the residues at or near both ends form a disulfide bond, giving peptide a loop-like configuration, which can be utilised to enhance immunogenic properties of the peptides. Protein is capable of eliciting, and/or binding with, neutralising antibodies. The neutralising domain is bounded by cysteine residues which occur at positions 296 and 331. The peptides can be used as immunogens or screening reagents to generate or identify poly- or monoclonal antibodies. See also ARR04427-R04506 and AAQ04273-Q04279. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hydrophilic; antigenic determinant; HIV; envelope; glycoprotein; env; gp recognition; B lymphocyte; type specific; antibody; vaccine; protection; immune response; infection; neutralisation; epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Revised record issued on 09-SEP-2004 : Correction to Feature Table Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                          Principal neutralising domain of HIV variants - used for producing peptide(s) and antibodies for diagnosis, prophylaxis and/or therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 68; DB 2; Length 25;
Pred. No. 0.0032;
1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                Claim 8 (58); Page 76; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW32887 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                              therapy therapy of HIV infection.
                                                                                                                                         88US-00252949.
89US-00359543.
89US-00407663.
                                                                                                         88US-00252949
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86.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-JAN-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                   JR, Putney SD,
J, Petrobre J;
                                                                                                                                                                                                                 (REPK ) REPLIGEN CORP.
                                                                                                                                                                                                                                                                                                          WPI; 1990-147824/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 25 AA;
                                                                                                                                        03-OCT-1988;
01-JUN-1989;
19-SEP-1989;
                                                                                                      03-OCT-1988;
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                                WO9003984-A
                                                                    19-APR-1990
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Synthetic.
                                                                                                                                                                                                                                                                      Lynn DU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                 ö
                   The peptides of the invention comprise the beta-turn AA SQ GFG or GPGR at postitions 312-315 in the AA numbering of HILV-IIIB (BH10) and flanking AA SQB having a length equal to or greater than 1 and pref. equal to or greater than 2 AAs; variants in which the GFG or GPGR SQ has been replaced by a different beta-turn SQ; and variants in which the FNH2-terminal gp AA and/or the free carboxyl terminal gp. AA has been blocked or modified otherwise. (Updated on 27-AUG-2003 to correct OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      T-cell, epitope, HIV-1, core protein, p24E, B-cell; antigen, gp160, gag, pol; vaccine, multimeric peptide, AIDS, 3D organisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel tandem synthetic HIV-1 peptide(s) - comprising T-cell epitope of gag protein linked to B-cell epitope of V3 loop protein of an HIV-I
                                                                                                                                                                                                                                                                                                                                 Gарв
                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                Query Match 87.0%; Score 67; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 0.0033; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 16; 69pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR68680 standard; peptide; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B cell epitope, LAI.
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                                                                                                                                                                                                                                      Sequence 17 AA;
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07-SEP-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tandem synthetic HIV peptide(s) useful as immunogens - comprising gag
protein T-cell epitope linked to env protein B-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                      HIV; human immunodeficiency virus; gag; T-cell; B-cell; epitope; env; V3 loop; vaccine; determinant; chimaeric.
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Pred. No. 0.0038;
0; Mismatches 1; Indel8
                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                               HIV-1 strain IIIB env protein V3 loop B-cell epitope.
Score 67; DB 2;
Pred. No. 0.0038;
0; Mismatches 1
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92.9%;
    87.0%;
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                                                                                                                            7 RIORGPGRAFYTIG 20
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                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
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                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-332082/30.
                     Local Similarity
es 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-JUN-1993;
                                                                                                                                                                                                                                                                                                                        25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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                                                                                                                                                                                                                                                                              AAW25898;
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Best Local S
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                                                                                                                                                                                              RESULT 160
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Matches
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ID AAW2
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AC AAW2
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The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-cell epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitopes are generally designed based on the p24 core protein and the B-cell epitopes from the V3 loop of the gp120 protein from various HIV-1 strains. This peptide corresponds to the V3 loop peptide epitope from the HIV-1 strain LAI. The peptide is used to generate a hybrid T- and B-cell epitope (AAW67353). (Updated on 17-OCT-2003 to standardise OS field)
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                                                                                                                                                                                                                                       Synthetic human immunodeficiency virus-1 peptide(s) - containing T-cell epitope and B-cell epitope(s) are candidate vaccines against HIV-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic chimeric HIV polypeptides - comprising gag protein T-cell epitope linked to gp41 B-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV-1; human immunodeficiency virus; vaccine; T-cell epitope;
gag protein; B-cell epitope; gp41 protein; chimeric; infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 67; DB 2; Length 20;
Pred. No. 0.0038;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
                                                                                                                                                  Sia CDY;
                                 93US-00073378.
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94US-00257528.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CONN-) CONNAUGHT LAB LTD.
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Best Local Similarity 92.9
Matches 13; Conservative
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                                                                                                                                                Chong P, Klein MH,
                                                                                                                                                                                             WPI; 1998-556461/47
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                                 09-JUN-1993;
09-JUN-1994;
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09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US5876731-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW99974;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW99974
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to new synthetic peptides comprising at least one amino acid sequence comprising an HIV gag protein T-cell epitope linked at its C- or N-terminus to an amino acid sequence comprising a B-cell epitope of the V3 loop of an HIV env protein, which can be used to generate vaccines against HIV-1. The T-cell epitope sequence is pref. P24M and P24H while the B-cell epitopes are derived from HIV strains including CTLB-56, V3MM, CTLB-29, CTLB-55, SF2, LAI, IIIB, RF, Z6, 2054, 1714 and BX08. The peptides are chimaeric and can be linked to a branched by backbone. This sequence represents the HIV-1 strain LAI env protein V3 loop B-cell epitope. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunogen; vaccine; HIV-1; T-cell; B-cell; epitope; core protein; gp120;
                                                                                                               HIV; human immunodeficiency virus; gag; T-cell; B-cell; epitope; env; V3 loop; vaccine; determinant; chimaeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tandem synthetic HIV peptide(s) useful as immunogens - comprising gag protein T-cell epitope linked to env protein B-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             87.0%; Score 67; DB 2; Length 20; 92.9%; Pred. No. 0.0038; live 0; Mismatches 1; Indels
                                                                    HIV-1 strain LAI env protein V3 loop B-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV-1 strain LAI V3 loop peptide epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW67366 standard; peptide; 20 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 8; Col 74; 41pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                        94US-00257528
                                                                                                                                                                                                                                                                                                                                                                  93US-00073378
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chong P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95US-00464329
                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RIQRGPGRAFYTIG 20
                                                                                                                                                                                                                                                                                                                                                                                                             (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
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       (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Klein MH, Sia CDY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-332082/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 AA;
  25-MAR-2003
20-OCT-1997
                                                                                                                                                                                                                                                                                                                   09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                  09-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUN-1995;
                                                                                                                                                                                                                                US5639854-A
                                                                                                                                                                                                                                                                        17-JUN-1997
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25-JAN-1999
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                                                                                                                                                                                   Synthetic.
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1D AAW6

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DT 17-C

DT 25-J

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Gaps

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This sequence represents a fragment of a HIV1 protein, and can be used in the immunogenic composition of the invention. The composition comprises a synthetic fusion polypeptide which includes a sequence encoding 1 or more T cell epitopes and a sequence encoding 1 or more E cell epitopes and a carrier. Both the T cell and B cell epitopes are derived from HIV proteins. The composition are useful as vaccines against HIV infection. The composition induces HIV1-specific polyclonal antibodies that are opponising and antiviral. The peptide components may be selected to induce a response against different viral isolates and in subjects who recognise different T cell epitopes. (Updated on 17-OCT-2003 to
its C terminus to an amino acid sequence containing a B-cell epitope of an HIV gp41 protein and containing the amino acid sequence: XILKDWX2; where X1 = E, A, G or Q, and X2 = A or T, or an amino acid sequence appable of eliciting an HIV-specific antiserum and recognizing the sequence XILKDWX2. The synthetic peptide is useful in vaccines against HIV infection and in diagnostic applications. AAW98892 to AAW98906, and AAW99899 to AAW99989 represent synthetic peptides from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency virus, useful antigens in producing vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunogenic composition; T cell epitope; B cell epitope;
                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                       ö
                                                                                                                                                                                      Length 20
                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                  Score 67; DB 2;
Pred. No. 0.0038;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                       AAY39699 standard; peptide; 20 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure, Col 9, 43pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            infection; antibody; antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Klein MH, Chong P, Sia CDY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93US-00073378
94US-00257528
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                                                                                                                                                                                    87.0%;
92.9%;
                                                                                                                                                                                                                                                      1 RIQRGPGRAFVTIG 14
                                                                                                                                                                                                                                                                       7 RIORGPGRAFYTIG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV1 chimeric peptide LAI
                                                                                                                                                                                                Local Similary
hes 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-550482/46.
                                                                                                                                                     Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-JUN-1995;
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26-NOV-1999
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                                                                                                                       invention
                                                                                                                                                                                                                                                                                                                                                                                                      AAY39699;
                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                      RESULT 164
                                                                                                                                                                                                                       Matches
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Length 20;

Score 67; DB 2; Pred. No. 0.0038;

87.0%; 92.9%;

Query Match Best Local Similarity

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The present sequence comprises a glycosylated fragment of gp120.

Glycopeptides comprising a fragment of the MUC1 repeat sequence,
CG especially having a Gal-GalNac or GalNac moiety on Thr-10 or Thr-11 (see
CAP96172-74), are useful as immunomodulators, causing super-proliferation
CG T cells. Such glycopeptides can be used in the treatment or diagnosis
CG a disease, in particular cancer, or as vaccine adjuvants. The
CJYCopeptides are particularly useful in manufacturing a medicament for
preventing or treating cancer by stimulating T cells whose receptors
CT recognize the glycopeptide. They are also useful for diagnosing or
CT recating cancer, e.g. carcinoma (e.g. mammary, lung, bladder or colon
Carcinomas, or ovary and endometrial tumours), or asrcoma (e.g. soft
Lissue and bone asrcomas, or leukaemia). Human peripheral blood
Lymphocytes were used in a proliferation assay. The proliferation index
Cf the gp120 glycopeptide (taking the index as 1 when no glycopeptide was
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                                                                                                                                                                                                                                                                               gpl20; MUC1; immunomodulator; glycopeptide; T-lymphocyte; T-cell;
proliferation; cancer; sarcoma; carcinoma; leukaemia; diagnosis; therapy;
vaccine; adjuvant; glycosylation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New immunomodulating glycopeptide that causes super-proliferation of 1 cells, useful for treating cells in vitro, for diagnosing or treating cancer (e.g. carcinoma or sarcoma) or as an adjuvant.
 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gape
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                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "O-glycosylated by GalNAc-beta-1-3Gal"
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 Indels
 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
 Mismatches
                                                                                                                                                                                                                                                    31cyoprotein gp120 glycosylated peptide.
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                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 24; 35pp; English.
                                                                                                                                               AAY96191 standard; peptide; 18 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Taylor-Papadimitrion
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99GB-00004695
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 RIORRGPGRAFVTICK 18
                                  1 RIQRGPGRAFVTIG 14
                                                                RIORGPGRAFYTIG 20
                                                                                                                                                                                                                   (first entry)
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 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 AA;
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                                                                                                                                                                                                                                                                                                                                                                                    Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                   Unidentified
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                                                                                                                                                                                                                   19-DEC-2000
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                                                                                                                                                                                  AAY96191;
 Matches
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HIV-1 strain IIIB gp120 variable region 3 epitope.

(first entry)

(revised) (revised)

25-MAR-2003 30-MAR-1998 17-0CT-2003

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New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IGH) protein with a mammalian, especially human, IGH chain fused in frame at its N-terminus to one or more human immune deficiancy virus (HIV) gpl20 epitopes. Fusion IG proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, as prevents or inhibits 'bystander' apoptoeis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral T cell immunity
                                                                                                                                                                             B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                          Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     85.7%; Score 66; DB 2; Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 10; Page 119; 154pp; English
                                        AAW76863 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (AMNA-) AMERICAN NAT RED CROSS.
                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US002766.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0040581P
                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zambidis E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-506315/43.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 14 AA;
                                                                                                                                                                                                                                                                                        Mus sp.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                          WO9836087-A1
                                                                                                                                                                                                                                                                                                                                                                                                               13-FEB-1998;
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                                                                                                           25-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                              20-AUG-1998.
                                                                        AAW76863;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scott D,
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This sequence represents an epitope corresponding to the central region of the variable region from the gpl20 protein of the human immunodeficiency virus type 1 (HIV-1) strain IIIB. The peptides AAW361S1-W361S6 were used to generate the multiple epitope-containing polypeptides to a mutant version of the first 47 amino acids of the P64k protein of Neisseria meningitidis B:4:P1.5 (AAW36149). The Neisseria peptide is used as a stabilisation peptide for purification of the heterologous protein, especially by immunoaffinity chromatography, when the fusion protein is produced in E. coli. (Updated on 2-MAR-2003 to correct PI field.) (Updated on 17-0CT-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Fusion protein for use as immunogen in vaccines - contains stabilising peptide derived from N-terminal 47 amino acids of Nisseria meningitidis
                                                                                           Mutant, P64k; fusion protein; stabilisation peptide; purification; immunosffinity chromatography; epitope; vaccine; human; animal; HIV-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gapa
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                                                                                                                                                                                                                                                                                                                                                       CE;
                                                                                                                                                                                                                                                                                                                                                    Gomez Rodriguez CE, Leal Angulo MDJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  85.7%; Score 66; DB 2; Length 15; 100.0%; Pred. No. 0.0041; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                      Alvarez Acosta A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV-1 isolate IIIB gp120 V3 loop epitope peptide.
                                                                                                                                                                                                                                                                                                     (INGG-) CENT ING GENETICA & BIOTECNOLOGIA.
                                                                                                                                                                                                                                                                                                                                 Duarte Cano CA, Guillen Nieto GE, Alva:
Carpio Munoz EL, Quintana Vazquez D, G
Silva Rodrigez RDLC, Nazabal Galvez C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; Page 12; 49pp; Spanish.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAM49356 standard; peptide; 15 AA.
                                                                                                                                            Human immunodeficiency virus 1,
                                                                                                                                                                                                                                         97WO-CU000001.
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                                                                                                                                                                                                                                                                                                                                                                                   Martin Dunn AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P64k antigen.
                                                                                                                                                                                                                                         17-JAN-1997;
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                                                                                                                                                                                                         24-JUL-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAM49356;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAM49356
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Gaps

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0.0039; haa 0; Indels

100.0%; Pred. No. 0.0 ive 0; Mismatches

Best Loca Matches

AAW36156 standard; peptide; 15 AA.

RESULT 167 AAW36156 AAW36156;

2 RIORGPGRAFVTI 14 1 RIORGPGRAFVTI 13 13; Conservative

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The invention relates to a composition comprising: (a) a first nucleic acid sequence that is a truncated A subunit coding region obtained or derived from a bacterial ADP-ribosylating exotoxin; and (b) a second nucleic acid sequence that is a truncated B subunit coding region obtained or derived from a bacterial ADP-ribosylating exotoxin. Each of the truncated subunit coding regions has a 5' deletion and encodes a subunit peptide not having an amino terminal bacterial signal peptide. The composition is useful for eliciting an immune response against an antigen or for manufacturing a medicament for enhancing an immune cresponse in a vertebrate subject (specifically a human) against an antigen. The composition is particularly useful as nucleic acid adjuvants for use in immunisation techniques. The present sequence is that of a HIV Spl20 peptide, used in examples of the invention to test for the adjuvant effects of plasmids phyv2002 and pplv2003 in enhancing the humoral and cellular immune responses to HIV-1 gpl20. (Updated on 23-OCT-2003 to
                                                                                                                                                                                                                                                            New compositions comprising nucleic acid adjuvants, useful in
immunization techniques, particularly for eliciting or enhancing an
immune response against an antigen in a human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacterial exotoxin; immune response; bacterial infection; vaccine; Human immunodeficiency virus; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 66; DB 6; Length 15;
Pred. No. 0.0041;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                            Example 5; Page 70; 143pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH60862 standard; peptide; 15 AA.
                                                                                                                 (POWD-) POWDERJECT VACCINES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                    26-NOV-2001; 2001WO-US043151.
                                                                    27-NOV-2000; 2000US-00724315.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-NOV-2001; 2001US-00993307.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-NOV-2000; 2000US-0253381P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              85.7%;
86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIQRGPGRAFVITGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Local 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                 Haynes JR, Arrington JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Haynes JR, Arrington J;
                                                                                                                                                                                                               WPI; 2003-221541/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HAYN/) HAYNES J R. (ARRI/) ARRINGTON J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV gp120 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2003162733-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH60862;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADH60862
셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to the preparation of antigenic structures with a high cross-reactivity which synergistically enhance immune responses to systematically and/or mucosally administered peptide antigens. Dendrimeric cructures containing a desired epitope are synthesised and coupled to a carrier molecule, and are then mixed with a CD4 fusion domain peptide (or a conjugate containing it) and an adjuvan such as alumina. The antigenic cructures are used in prophylactic or therapeutic vaccine formulations for the prevention or treatment of infections, autoimmune diseases or cancer in humans and other animals, and may also be used as part of alganostic systems. The antigenic structures of the invention have synergistically enhanced cross-reactivity and immunogenicity compared with the parent dendrimeric epitope structures from which they are contained. Sequences AAM49354-AAM49359 represent gpl20 v3 loop epitopes from different HIV-1 isolates. These were used in an exemplification to demonstrate that the potentiation of immunogenicity and cross-reactivity associated with the structures of the invention is independent of the sequence of the v3 loop peptide used in the antigenic structure. The sequence of the v3 loop peptide used in the antigenic structure. The sequence represents the gpl20 v3 loop epitope from HIV-1 isolate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                   Preparation of antigenic structures having enhanced cross-reactivity and immunogenicity, useful for diagnosis or in vaccines, e.g. against cancer, comprising dendrimeric antigenic epitope conjugated with carrier
                                                                                                                                                                                                                                        Aguilar Rubido JC, Iglesias Perez E;
Garay Perez HE, Muzio Gonzalez VL, Guillen Nieto GE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP-ribosylating exotoxin; immune response; immunisation; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   85.7%; Score 66; DB 5; Length 15; 100.0%; Pred. No. 0.0041; Anismatches 0; Indels
                                                                                                                                                                                             (INGG-) CENT ING GENETICA & BIOTECNOLOGIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Fig 1c; 41pp; Spanish.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR39121 standard; peptide; 15 AA.
                                                                                                                                                                                                                                        Cruz Ricondo LJ, Aguilar Rubido
Reyes Acosta O, Garay Perez HE,
Duarte Cano C, Penton Arias E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV-1 gp120 peptide SEQ ID NO 21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus 1.
                                                                                             01-NOV-2001; 2001WO-CU000007.
                                                                                                                                             03-NOV-2000; 2000CU-00000242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 RIQRGPGRAFVTI 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 RIORGPGRAFVTI 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            adjuvant; HIV; gp120
                                                                                                                                                                                                                                                                                                                                        WPI; 2002-444347/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003004055-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 AA;
WO200236160-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-JAN-2003
                                             10-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR39121;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Matches

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Gaps

WPI; 2003-897945/82.

immunodeficiency virus 1. (UYNY ) UNIV NEW YORK STATE. (YEDA ) YEDA RES & DEV CO LTD. 04-FEB-2004; 2004WO-US003304. 04-FEB-2003; 2003US-044682P WO2004069863-A2. Anglister J, 19-AUG-2004 Synthetic. ADR18865; peptide. Human RESULT 171 ઠે g 

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                                                                                                                                                                                                                                      The present invention relates to a new composition comprising first and second nucleic acid sequences. The first or second nucleic acid sequence is a truncated A or B subunit coding region, respectively, derived from a bacterial ADP-ribosylating exctoxin, with the provision that each of the truncated subunit coding regions has a 5' deletion and encodes a subunit peptide not having an amino terminal bacterial signal peptide. The invention is useful for enhancing an immune response against bacterial infection. The present sequence is Human immunodeficiency virus gp120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein co-ordinate data; V3 loop peptide; HIV-1; envelope glycoprotein; gp120; human monoclonal antibody 447-52D; murne monoclonal antibody 0.5 beta; immunogen; immunogenic; V3 loop epitope; HIV-1 infectivity inhibitor; anti-HIV; vaccine; HIV-1 infection.
New composition comprising first or second nucleic acid sequence, which is a truncated A or B subunit coding region obtained or derived from a bacterial ADP-ribosylating exotoxin, useful as a vaccine against bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 66; DB 7; Length 15;
Pred. No. 0.0041;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        V3-IIIB beta-hairpin related peptide SEQ ID NO:24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   three-dimensional atomic structural conformation;
                                                                                                                                                                        Example 5; SEQ ID NO 21; 72pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RIORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIORGPGRAFVITCK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 86.7
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 AA;
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cc isolated peptide molecule or isostere that mimics the three-dimensional CC isolated peptide molecule or isostere that mimics the three-dimensional cc envelope glycoprotein gp10 that is bound to, and constrained by, human monoclonal antibody (Mab) 447-20, murine MAD 0.5 beta or an antigen convocational antibody (Mab) 447-20, murine MAD 0.5 beta or an antigen conformation from the same V3 loop peptide when it is in free differs in conformation from the same V3 loop peptide when it is in free compounds a molecule that is useful as an HV10 13 loop jemunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor con the surface of a receptor-bearing target cell; (3) a composition (C2) that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor con the surface of a receptor-bearing target cell; (3) a composition (C3) con the surface of a receptor-bearing target cell; (3) a composition (C3) con the surface of a receptor-bearing target cell; (3) a composition (C3) con the surface of a receptor-bearing target cell; (3) a composition (C3) con the surface of a receptor-bearing target cell; (3) a composition (C3) con the surface of a receptor-bearing target cell; (3) a composition (C3) con the surface of a receptor-bearing target cell; (3) a composition (C3) con a receptor-administration of an anti-HIV-1 antibody response specific for a vectoper of a receptor-bearing target cell; (4) an interaction of HIV-1 with an R5 conformational (C4) useful for blocking the interaction of HIV-1 with a mit conformationally constrained by NC; and conformationally constrained structure of a value of a constrained HIV v3 view peptide; (7) a computer generating (C3) and computer readable medium (C4) computing phase of a constrained HIV v3 view peptide; (7) a computer generating conformationally constrained by being bound to 44-520 or 0.5beta MAD or its antigen binding fragments, comprising a 3D atomic structure of a v3 loop peptide that is conform

neutralising antibody response specific for a V3 loop epitope. (C4) is useful for preventing an HIV-1 infection in an uninfected subject at risk for such infection or for inhibiting viral spread and disease progression

sequence represents a peptide used in

an infected subject. The present sequence exemplification of the present invention.

Sequence 16 AA;

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                                                                                                                                                                                                   Human immunodeficiency virus; retrovirus; vaccine; antibodies; HbC; HBe; antigen; hepatitis B virus; HBV; core.
                       Gaps
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85.7%; Score 66; DB 8; Length 16; 100.0%; Pred. No. 0.0044; ive 0; Mismatches 0; Indels
                                                                                                                                                                                Epitope comprising residues 308-327 of HIV env gp 120.
        100.0%; Pred. ...
                                                                                                           AAR04060 standard; peptide; 21 AA.
                                                                                                                                                                                                                                                                                            88JP-00220770.
                                                                                                                                                    (revised)
(first entry)
                                        1 RIORGPGRAFVTI 13
                                                     4 RIQRGPGRAFVTI 16
         Local Similarity 100.
                                                                                                                                                                                                                                                   JP02069194-A.
                                                                                                                                                                                                                                                                                           02-SEP-1988;
                                                                                                                                                   25-MAR-2003
                                                                                                                                                             23-JUL-1992
                                                                                                                                                                                                                                                                       08-MAR-1990.
                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                AAR04060;
  Query Match
                                                                                        RESULT 172
            Best Loc
Matches
                                                                                                   AAR04060
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Composition for inhibiting HIV-1 infection, comprises isolated peptide molecule that mimics atomic structural conformation of V3 loop peptide of HIV-1 envelope glycoprotein that is bound to, and constrained by human monoclonal antibody.

Rosen O;

Sharon M, Schapira M, Zolla-Pazner S,

WPI; 2004-625447/60.

The present invention describes a composition (C1) which comprises an

Example 7; SEQ ID NO 24; 127pp; English.

us-08-869-386-1.rag

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type 1 (HIV-1) V3IIIB peptide used in the exemplification of the invention. (Updated on 29-AUG-2003 to standardise OS field)
                                                             85.7%;
                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13 RIQRGPGRAFVTI 25
                                                                                                             1 RIORGPGRAFVTI 13
                                                                                                                                   12 RIORGPGRAFVTI 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIORGPGRAFVTI 13
                                                                                      13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         against HIV infections.
                                                                                                                                                                                                                                                (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-341488/42.
                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 25 AA;
                                      Sequence 24 AA;
                                                                                                                                                                                                                                                16-OCT-2003
25-MAR-2003
29-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                       15-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                       WO9423746-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-APR-1993;
                                                                                                                                                                                                                         AAR63820;
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                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW22329
ID AAW2
XX
                                                                                                                                                                        RESULT 17
AAR63820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a composition comprising activated alpha-2-. macroglobulin (alpha_2M asterisk ), 3-O-deacylated monophosphoryl lipid A (MPL) and granulocyte macrophage colony stimulating factor (GM-CSF). The invention also relates to an adjuvant suitable for use in multivalent HIV immunogenic compositions. The compositions is useful for eliciting an immune response. The present sequence is human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Use of 2-macroglobulin (2Masterisk), 3-O-deacylated monophosphoryl lipid A (MPL) and granulocyte macrophage colony stimulating factor (GM-CSF) for eliciting an immune response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus type 1; HIV-1; adjuvant; immunomodulator; alpha-2-macroglobulin; 3-0-deacylated monophosphoryl lipidA; MPL; GM-CSF; granulocyte macrophage colony stimulating factor; immune response; vaccine; V3IIIB peptide.
                                                                                                                                          The synthetic epitope is used in a complex with either the hepatitis B core antigen (HBc) or a sol. cleavage prod. of HBc (HBe), to prepare a vaccine. The peptide corresponds to residues 308-327 of the HIV env glycoprotein 120, with an N-terminal initiation Met. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                           Gaps
                                                                                       Antigen granule comprising HBC or HBE antigen - and HIV neutralised epitope obtd. by expression of recombinant prod., for e.g. vaccine.
                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus type 1 (HIV-1) V3IIIB peptide.
                                                                                                                                                                                                                                                 Length 21;
                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                85.7%; Score 66; DB 2; I
100.0%; Pred. No. 0.0056;
                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                           Claim Disclosure; Fig 4; 11pp; Japanese.
                           (KAGA ) KAGAKU OYOBI KESSEI RYOHO
                                                                                                                                                                                                                                                                                                                                                                                       AAE20149 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 21; 53pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patel DD;
   88JP-00220770.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-AUG-2001; 2001WO-US026589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-AUG-2000; 2000US-0227624P
                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                  1 RIQRGPGRAFVTI 13
                                                                                                                                                                                                                                                                                                                  RIORGEGRAFUTI 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Liao H,
                                                   WPI; 1990-119518/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-269315/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYDU-) UNIV DUKE
                                                               N-PSDB; AAQ02417.
                                                                                                                                                                                                                          Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200215930-A1
   02-SEP-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                        29-AUG-2003
18-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Haynes BF,
                                                                                                                                                                                                                                                                                                                                                                                                                AAE20149;
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AAR63809-R63849 are epitopes from the human immunodificiency virus type I (HIV-1) gp120, by binding one or more of these epitopes to a carrier a HIV vaccine is produced. These vaccines can elicit the production of HIV-neutralishing antiboddes in monkeys, and therefore may be used to prevent HIV infections, and to highten the immune response in HIV infected humans. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New peptide(s) comprising HIV gpl20 epitope(s) - for prodn. of vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus type 1; HIV-1; gp120 epitopes; vaccines; HIV neutralising antibodies.
                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                            ô
85.7%; Score 66; DB 5; Length 24; 100.0%; Pred. No. 0.0063; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 66; DB 2; Length 25;
Pred. No. 0.0065;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Horal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jeansson S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85.7%; Scor.
100.0%; Pred. No. v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV-1 gp120-23 epitope amino acida 296-230.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vahlne A, Svennerholm B, Rymo L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW22329 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                           AAR63820 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 18; 77pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SYNT-) SYNTELLO VACCINE DEV AB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94WO-SE000340.
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The invention relates to a novel monoclonal antibody (MAb) NM03 which binds to epitopes from the human immunodeficiency virus type 1 (HIV-1). The antibody was raised conventionally by immunising Balb/c mice with purified live HV-1 MM, then isolating splenocytes and fusing them to P3- X63-Ag8-U1 cells. Hybridomas were then screened with membrane fractions from infected H9 cells. The MAb was observed to bind to a protein band of 120 kD on a Western blot of separated, denatured HIV-1 proteins. This binding was shown to be between residues 320-327 by copteins. This binding was shown to be between residues 320-327 by colinibit infection of cells by HIV-1 shown by infecting H9 cells with live strains of HIV-1 and testing infection by a p24 assay. This peptide sequence represents the V31 loop region from HIV-1 clinical strain 9622, where the MAb NNO3 binds. The MAb can be used for the passive immunisation of humans susceptible to, or infected with HIV-1. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                        Epitope; human immunodeficiency virus type 1; HIV-1; gpl20; gpl60; monoclonal antibody; V3 loop; immunisation; mouse; splenocyte; hybridoma; membrane fraction; passive immunisation; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monoclonal antibody specific for human immunodeficiency virus type 1 MN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 64; DB 2; Length 19;
Pred. No. 0.01;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 strain - for passive immunisation against infection.
                                                                                      HIV-1 clinical strain 9622 gp120 V3 loop peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide sequence of the specification.
                                                                                                                                                                                                                                                                                                                                                                       (NISP ) NISSIN SHOKUHIN KAISHA LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW62892 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Col 10; 14pp; English.
                                                                                                                                                                                              Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                     94US-00279906
                                                                                                                                                                                                                                                                                                                                                                                                         Yoneda Y, Ohno T, Terada M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   83.1%;
                                                                                                                                                                                                                                                                                                                                       94US-00279906
                                      (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 ORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGPGRTFVTIGK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 92.3
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-225475/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 AA;
                                                                                                                                                                                                                                                                                                 25-JUL-1994;
                                                                                                                                                                                                                                                                                                                                      25-JUL-1994;
                                    17-OCT-2003
18-SEP-1997
                                                                                                                                                                                                                               US5618922-A
                                                                                                                                                                                                                                                                  08-APR-1997.
   AAW22329;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW62892
IID AAW6
XX
AC AAW6
XX
XX
DT 30-S
XX
XX
KW MONO
XX
KW HIV
XX
OS Synt
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à a

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AAW62889-900 represent peptides used to identify a peptide sequence (AAW62874) present in HIV-1gp120 or gp160 which is bound by the monoclonal antibody of the invention. The antibody neutralises in vitro the infection of H9 cell by an active HIV strain MN according to the p24 analytical method. The antibody is used for treatment of human HIV
                                                                                                                                                                                                                                                                                                                                                                            Monoclonal antibody which binds to HIV-1gp120 or gp160 - used to prevent and treat human HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Methods for treating immunoinfective cluster virus infections - utilise antibodies or fragments characteristic of auto antibodies produced by patients with rheumatic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         autoantibody; immunoinfective cluster virus; nuclear protein antigen; systemic rheumatic disorder; human immunodeficiency virus; HIV-1; systemic lupus erythematosus; mixed connective tissue disease; scleroderma; glycoprotein 120; glycoprotein 41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 2; Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Small ribonucleoprotein complex; U1 snRNP; 70K protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1 gpl20/41 protein consensus binding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 64; DB 2;
Pred. No. 0.01;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                        (NISP ) NISSIN SHOKUHIN KAISHA LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR62152 standard; peptide; 12 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Page 8; 12pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ehresmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYSC-) UNIV SOUTHERN CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1,
                                                                                                                                                                                                    96JP-00344904.
                                                                                                                                             96JP-00344904
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           83.1%;
92.3%;
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Best Local Similarity 92.3.
Local 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
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                                                                                                                                                                                                                                                                                                                  WPI; 1998-433774/37.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 AA;
                        JP10182489-A.
                                                                                                                                         25-DEC-1996;
                                                                                                                                                                                                    25-DEC-1996;
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                                                                               07-JUL-1998,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9420141-A1
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02-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infection
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(first entry)

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Glutamic acid; glutamine; vaccine; major histocompatibility complex; MHC; immunomodulator; antiallergic; endocrine; neuroprotectant; virucidal; bactericidal; antiparasitic; fungicidal; cytostatic; medicine; pharmaceutical; immune disorder; immune deficiency; autoimmune; hypersenaitivity; allergy; graft rejection; infection; hormonal disorder; central nervous system disease; cancer; melanoma; anti-melanoma vaccine; human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stabilized pharmaceutical containing N-terminal glutamic acid or glutamine, useful e.g. in anti-melanoma vaccines, is an addition salt
                                                                                                                                                    Vaccine related MHC ligand peptide SEQ ID NO:535.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Klinguer-Hamour C, Corvaia N, Beck A,
                                                                                                                                                                                                                                                                                                                                                                                                                                      (FABR ) FABRE MEDICAMENT SA PIERRE.
                                                                  AAM99432 standard; peptide; 12 AA.
                                                                                                                                                                                                                                                                                                                                                                                                          23-MAR-2000; 2000FR-00003711.
                                                                                                                                                                                                                                                                                                                                                                              22-MAR-2001; 2001WO-FR000872.
1 RGPGRAFVTIGK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-611470/70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  with strong acid.
                                                                                                                                                                                                                                                                                                                         WO200170772-A2
                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                          07-DEC-2001
                                                                                                                                                                                                                                                                                                                                                    27-SEP-2001.
                                                                                                AAM99432;
                                                                                              The UI snRNP is the target of high-titre, high avidity autoantibodies occurring in the systemic rheumatoid disorders of mixed connective tissue disease, scleroderma and systemic lupus erythematosus. It has been found that some sites in the UI snRNP 70K protein (see AAR62120-R62183) are homologous to sites in HIV-1 gal20/41 (AAR62136-R62182) and that anti-RNP autoantibodies can be used to neutralise HIV-1. In particular, the sequence AAR62152 from HIV-1 gpl20 matches a consensus binding sequence which is necessary and sufficient for high affinity binding to UI RNA. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG-2003 to correct OS field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptides AAW54903-W54941 can be used as an immunoadsorbent in an immunoassay for detecting antibodies to HIV gpl20, or as an immunogen for eliciting antibodies to HIV in a mammal
                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunoadsorbent; immunoassay; HIV gp120; immunogen; antibody; Human.
                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gp120 peptides - useful as immunoassay reagents or vaccine
                                                                                                                                                                                                                       81.8%; Score 63; DB 2; Length 12; 100.0%; Pred. No. 0.0096; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 81.8%; Score 63; DB 2; Length 12; Best Local Similarity 100.0%; Pred. No. 0.0096; Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV gp120 envelope protein, peptide 127, analogue 267d.
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              Disclosure; Page 56; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                AAW54932 standard; peptide; 12 AA.
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91US-00726605.
94US-00326676.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UNBI-) UNITED BIOMEDICAL INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95US-00488252.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88US-00155321.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-1998 (first entry)
                                                                                                                                                                                                                                                      12; Conservative
                                                                                                                                                                                                                                                                                 4 RGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                           1 RGPGRAFVTIGK 12
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                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                              Sequence 12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 12 AA;
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09-JUL-1991;
19-OCT-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            components.
                                                                                                                                                                                                                                                                                                                                                                                                             AAW54932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wang CY;
                                                                                                                                                                                                                                                      Matches
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The present invention describes a pharmaceutical compound (I) that

contains an N-terminal glutamic acid (Glu) or glutamine (Gln) residue in

the form of an addition salt with a strong, physiologically acceptable

acid (II). Also described are: (a) a pharmaceutical composition

containing at least one (I); (b) a vaccine containing at least one (I)

where this is a major histocompatibility complex (MFIC) ligand (Ia); (c) a

method for in vitro diagnosis of diseases associated with the presence of

(Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process

corporation (I). (I) has immunomodulator, endocrine, antiallargic,

neuroprotectant, virucidal, bactericidal, antiparasitic, fungicidal and

cycostatic activities. (I) are useful, in human or veterinary medicine,

containing at least, hypersensitivity, allergy, graft

rejection, infection, hormonal disorders and central nervous system

contained with interaction of: (I) viral, bacterial, parasitic or fungal

infections; or (ii) of cancers. A particular application is in anti-

melanoma vaccines. (I) are also useful for in vitro diagnosis of diseases

associated with interactions between MHC and (I), e.g. melanoma and thuman

immunodeficiancy virus infection. AMM98989 to AAM9952 represent invention

immunodeficiancy virus infection. Compounds from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     81.8%; Score 63; DB 4; Length 12;
100.0%; Pred. No. 0.0096;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Pred. No. 0.0
ive 0; Mismatches
Claim 9; Page 122; 149pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.
1es 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 ORGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ò
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Gaps

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4 RGPGRAFVTIGK 15

(revised) (revised)

AAR04476;

1 QRGPGRAFVTIG 12

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New fusion immunoglobulin heavy chain including gpi20 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immunos deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, are bound to gpl20 protein, maximising induction of protective antiviral real immunity.
                                                                                                                                                B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                               Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   81.8%; Score 63; DB 2; Length 14;
100.0%; Pred. No. 0.011;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 10; Page 119; 154pp; English.
               AAW76864 standard; peptide; 14 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         81.0.,
100.0%; Fil
                                                                                                                                                                                                                                                                                                                                                                                                                                           (AMNA-) AMERICAN NAT RED CROSS.
                                                                                                                                                                                                                                                                                                                                                                         98WO-US002766
                                                                                                                                                                                                                                                                                                                                                                                                           97US-0040581P.
                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scott D, Zambidis E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-506315/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14 AA;
                                                                                  25-JAN-1999
                                                                                                                                                                                                                                                        Mus sp.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                         13-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                           13-FEB-1997;
                                                                                                                                                                                                                                                                                                      WO9836087-A1
                                                                                                                                                                                                                                                                                                                                        20-AUG-1998.
                                                 AAW76864;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Peptide RP140 comprises segments of the Principal Neutralising Domain (envelope protein) from isolates HIV-RF and HIV-IIIB. The last Cys cassidue is added for the purpose of crosalinking to carrier proteins. Cysteine residues may be added, so that the residues at or near both ends form a disulfide bond, giving peptide a loop-like configuration, which can be utilised to enhance immunogenic properties of the peptides. Protein is capable of eliciting, and/or binding with, neutralising antibodies. The neutralising domain is bounded by cysteine residues which occur at positions 296 and 331. The peptides can be used as immunogens antibodies. See also AAR04427-R04506 and AAQ04273-Q04279. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                              HIV isolates HIV-IIIB and HIV-RF; hybrid peptide PR140; therapy; AIDS; principal neutralising domain; antibodies; diagnosis; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Revised record issued on 09-SEP-2004 : Correction to Feature Table Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                               Principal neutralising domain of HIV variants - used for producing peptide(s) and antibodies for diagnosis, prophylaxis and/or therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                             Grimaila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 63; DB 2; Length 23;
Pred. No. 0.017;
1; Mismatches 1; Indels
                                                                                   Human immunodeficiency virus hybrid peptide RP140.
                                                                                                                                                                                                                                                                                                                                                            Javaherian K, Farley J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tip of HIV-IIIB V3 loop peptide of gp120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 8 (59); Page 76; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW38249 standard; peptide; 16 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy therapy of HIV infection.
                                                                                                                                                                                                                                                                        88US-00252949.
89US-00359543.
89US-00407663.
                                                                                                                                                                                                                                               88US-00252949.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                81.8%;
85.7%;
                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 IQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-MAR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12; Conservative
                                                                                                                                                                                                                                                                                                                                                             Putney SD,
                                                                                                                                                                                                                                                                                                                                 (REPK ) REPLIGEN CORP
                                                                                                                                                                                                                                                                                                                                                            Rusche JR, Putney SD
Lynn DU, Petrobre J;
                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1990-147824/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 23 AA;
                          09-SEP-2004
25-MAR-2003
20-SEP-1990
                                                                                                                                                                                                                                               03-OCT-1988;
                                                                                                                                                                                                                                                                           03-OCT-1988;
                                                                                                                                                                                                                                                                                      01-JUN-1989;
                                                                                                                                                                                      WO9003984-A.
                                                                                                                                                                                                                  19-APR-1990
                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW38249;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Matches
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Gaps

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AAR04476 standard; protein; 23 AA.

RESULT 181

AAR04476 ID AAR0 XX

4 RGPGRAFVTIGK 15

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Length 14;

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Norton EK; Pincus SE,

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Vaccine comprises recombinant, attenuated pox-virus - use for vaccinating against viral infections such as rabies, hepatitis B, HIV, HSV, EBV, CMV,
                                                                                                                                                                                                                                                                                                                                                                                                                  cytotoxic T lymphocytes (CTL). Mice were unoculated with cells transformed with NVAC recombinants expressing the HIV-1 IIIB env gene. These mice generated CTL's and memory precursors of CTL's. The env gene. These mice generated CTL's and memory precursors of CTL's. The transformed which NVAC is derived overnight with the peptide sequences given to test specificity of CTL antigen receptor recognition of the V3 loop region of HIV IIIB gpl.O. ALVAC is derived from a canarypox virus and NVAC is derived fron Copenhafen strain vaccinia virus which have been modified by deletion of non-essential regions of the genome encoding known or potential virulence factors. The deletion loci were engineered as recipient loci for the insertion of foreign genes. See also AAQ35501-864. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                         The sequences given in AAR27465-67 were used to perform assays for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81.2%; Score 62.5; DB 2; Length 21; 93.3%; Pred. No. 0.019; Live 0; Mismatches 0; Indels
                                                                                                                                                            Tartaglia J, No
J, Johnson GP,
                                                                                                                                                            Taylor J, Ta
Limbach KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus; V3 loop.
                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 295; 456pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cox WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR31219 standard; peptide; 21 AA
                                      91US-00666056.
91US-00713967.
92US-00847951.
  92WO-US001906.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91US-00715921.
92US-00897382.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RIORGPGRAFVT-GK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               V3 peptide from HIV-1 IIIB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tartaglia J,
                                                                                                                                                            , Perkus ME, 1
De Taisne C,
Gettig RR;
                                                                                                                        (VIRO-) VIROGENETICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (VIRO-) VIROGENETICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(revised)
                                                                                                                                                                                                                                          WPI; 1992-331718/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 AA;
  09-MAR-1992;
                                        07-MAR-1991;
11-JUN-1991;
                                                                              06-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9222641-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-OCT-2003
25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-DEC-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Paoletti E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-MAY-1993
                                                                                                                                                                               Riviere M,
Francis J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-JAN-2003
                                                                                                                                                              Paoletti B,
                                                                                                                                                                                                                                                                                                                               mumps etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR31219;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 184
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  셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ï
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence was used in the development of a novel multivalent chimeric peptide, comprising at least 2 tandemly repeated units, where the lift portion of the repeated unit comprises a human mucin il (MUCI) sequence which forms an extended connector and a base of a loop structure of human MUCI, and a 2nd portion comprising an immunoglobulin super family protein Omega loop sequence. In the peptide, the natural structure of MUCI tandem repeats can be used to present an Omega loop sequence in a functional conformation that is both multivalent and biologically active. It can provide prophylactic and therapeutic agents which have the binding specificity of an immunoglobulin super family member protein but do not have the entire protein's backbone. It can be used to neutralise pathogenic viruses, e.g. human immunodeficiency virus (HIV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cytotoxic T lymphocyte; CTL; NYVAC; ALVAC; HIV-1 IIIB; env; memory precursor; CTL antigen receptor; V3 loop; gp120; canarypox virus; Copenhagen strain; vaccinia virus; virulence factor; deletion loci;
                                                                                                                                                                                                                                                                                                                                                                                                                            New multivalent chimeric peptide(s) for neutralising pathogenic microbes - comprising a loop structure of human mucin 1 and an omega loop of an immunoglobulin superfamily protein.
Multivalent chimeric peptide; tandem repeat unit; human; mucin 1; MUC1; Omega loop sequence; prophylaxis; therapy; V3 loop peptide; pathogenic virus neutralisation; gpl20; HIV-IIIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ij
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81.2%; Score 62.5; DB 2; Length 16; 93.3%; Pred. No. 0.015; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sxample 2; Page 39; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR27465 standard; protein; 21 AA.
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0
                                                                                                                                                                                                                                                                                                          (POPU-) POPULATION COUNCIL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     V3 peptide from HIV-1 (IIIB).
                                                                                                                                                                                                                                             96US-00594403.
96US-00730244.
                                                                                  Human immunodeficiency virus
                                                                                                                                                                                                         97WO-US001726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 RIQRGPGRAFVT-GK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                Fontenot JD, Phillips DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-402551/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          recipient loci
                                                                                                                                                                                                      29-JAN-1997;
                                                                                                                                                                                                                                               31-JAN-1996;
15-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9215672-A1
                                                                                                                        409728187-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
24-FEB-1993
                                                                                                                                                                07-AUG-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR27465;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Loca
Matches
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%XCCCCCCCCCX8X4444X4X4X4X4X4X6XX8XX8X4X4X8X

AAR27465

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Gaps

1;

covalent linkages is comprised of maleimide derivatives; the (poly) anionic moiety is composed of one to five residues of the anionic (poly) anionic moiety is composed of one to five residues of the anionic profine of a carboxylic, sulphonic or phosphonic acid; the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b; and the PND peptide has a linear structure, a disulphide-bonded cyclic structure, an amide-bonded cyclic structure or a thioetherbonded cyclic structure. The present sequence (PND135-12) is an example of a PND peptide component used in the co-conjugate. The co-conjugate is useful for inducing anti-peptide immune response in mammals, for inducing HIV-neutralizing antibodies in mammals, for formulating vaccines to prevent HIV infection or disease, including AIDS, or for treating humans afflicted with HIV infection or disease

88666666666668888

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Gaps

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80.5%; Score 62; DB 2; Length 12; 100.0%; Pred. No. 0.014; ive 0; Mismatches 0; Indels

12; Conservative 1 RIORGPGRAFVT 12 RIORGPGRAFVT 12

Matches

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1;

Gaps

1;

Sequence 12 AA;

Query Match Local

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The peptide represents the V3 loop epitope of HIV-1 IIIB. The peptide was used to test the specificity of cytotoxic T lymphocyte antigen receptor recognition of the V3 loop region of HIV IIIB gpl20. See also AR31218-26. (Updated on 09-JAN-2003 to add missing OS field.) (Updated on 25-MAR-2003 to correct PN field.) (Updated on 24-OCT-2003 to standardise OS
                               Modified recombinant virus with inactivated non-essential genetic functions - comprises e.g. vaccinia or avipox virus, used as HIV vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "bonds to the OMPC of the conjugate via this site"
                                                                                                                                                                                                                                                                                                                                                                                                       AIDS; ARC; human immunodeficiency virus; vaccine; Neisseria; meningitidis b; outer membrane protein complex; OMPC; PND135-12.
                                                                                                                                                                                        Score 62.5; DB 2; Length 21;
Pred. No. 0.019;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                 Example 5; Page 65; 159pp; English
                                                                                                                                                                                                                                                                                                                                                                                                   AIDS; ARC; human immunodeficiency
                                                                                                                                                                                                                                                                                                             AAR31278 standard; peptide; 12 AA.
                                                                                                                                                                                                                                                                                                                                                                              HIV principal determinant peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90US-00555339.
90US-00555966.
91US-00715276.
91US-00715278.
                                                                                                                                                                                        81.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 91EP-00306598
                                                                                                                                                                                                                                   1 RIQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                12-FEB-1993 (first entry)
                                                                                                                                                                                     Query Match
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                 Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
Modified-site
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The invention relates to a conjugate of an HIV principal neutralizing determinant (PMD), or an immunologically equivalent peptide (PPE), covalently coupled to an immunogenic protein complex through an anionic polysaccharide linker. Pref. the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b and the PND peptide has a linear structure, a disulphide-bonded cyclic structure, an amide-bonded cyclic structure. HIV peptide-polysaccharide-protein conjugates - used in vaccines or to produce antibodies to prevent or treat HIV infection. 'note= "Joins onto polysaccharide-protein complex via Human immunodeficiency virus; principal neutralizing determinant; outer membrane protein complex; OMPC; Neisseria; AIDS; PND-135-12. HIV-PND-polyBaccharide-protein conjugate vaccine. Location/Qualifiers AAR26714 standard; peptide; 12 AA. ΕÀ Claim 9; Page 57; 63pp; English. 90US-00555558. 90US-00555974. 91US-00715275. 91US-00715277. Emini 90US-0055558 this site" (first entry) Tolman RL, MERI ) MERCK & CO INC. WPI; 1992-034437/05. Key Modified-site 19-JUL-1990; 09-FEB-1993 19-JUL-1990; 19-NUL-1991; 29-JAN-1992 EP468714-A. Marburg S, Synthetic AAR26714; RESULT 186 AAR26714 

The invention relates to a co-conjugate comprising an immunogenic protein or protein complex having a first set of covalent linkages to low molecular weight moieties which have an anionic or polyanionic character at physiological ph, and a second set of covalent linkages to peptides comprising HIV principal neutralizing determinants (PND's) or immunologically equivalent peptides. Preferably at least one set of the

Conjugate proteins comprising HIV peptide components - useful for preparing vaccines for e.g. AIDS or for treating infections.

Claim 12; Page 56; 63pp; English.

EA: Emini

Tolman RL,

Leanza WJ, Marburg S,

WPI; 1992-026505/04.

& CO INC.

(MERI ) MERCK

19-JUN-1991;

us-08-869-386-1.rag

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AAR58601 standard; peptide; 13 AA.
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              RESULT 188
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The present sequence (PND135-12) is an example of a PND peptide component. The conjugates are used for inducing HIV-neutralising antibodies or for making vaccines to prevent contraction of HIV infection of disease. The antibodies can be used for passively protecting against infection by HIV, or for protecting against proliferation of HIV postinfection, or for treating AIDS, or in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAREBGEO3 is an amino acid moiety (AAM), when inserted between amino acid residues 168-169 of alkaline phosphatase (AP) (AARSBEO8) it enables the enzymatic activity of AP to be modulated upon the binding of a molecule to the inserted sequence. The modified AP can now be used as a diagnostic assay, where the presence of specific antigens and antibodies can be inferred by changes in the enzymes activity. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genetically engineered hybrid enzymes and ligand conjugates - useful for diagnostic assay, to detect antigens and antibodies by changes in enzyme
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                                                                                                 80.5%; Score 62; DB 2; Length 12;
100.0%; Pred. No. 0.014;
rative 0; Mismatches 0; Indels
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100.0%; Pred. No. 0.015;
ive 0; Mismatches 0;
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Yue VT;
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Russell JC,
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les 12; Conser
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01-MAY-1995
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AARSB601 is an amino acid moiety (AAM), when inserted between amino acid residues 407-408 of alkaline phosphatase (AP) (AARSB608) it enables the enzymatic activity of AP to be modulated upon the binding of a molecule to the inserted sequence. The modified AP can now be used as a diagnostic assay, where the presence of specific antigens and antibodies can be inferred by changes in the enzyme's activity. (Updated on 25-MAR-2003 to correct PN field.)
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100.0%; Pred. No. 0.015;
tive 0; Mismatches 0
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                                                           Alkaline phosphatase 407-408 insertion.
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Yue VT;
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93US-00100708.
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Russell JC,
  (revised)
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hes 12; Conserv
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25-MAR-2003
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100.0%; Pred. No. v.
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Yue VT;
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Yue VT;
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March SC, Russell JC,
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                                                               AARS8605 is an amino acid moiety (AAM), when it replaces the amino acid residues 91-93 of alkaline phosphatase (AP) (AARS8608) it enables the enzymatic activity of AP to be modulated upon the binding of a molecule to the inserted sequence. The modified AP can now be used as a diagnostic assay, where the presence of specific antigens and antibodies can be inferred by changes in the enzymes activity. (Updated on 25-MAR-2003 to
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Cys(15) via an o-xylylene bridge, forming cyclic peptide"
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diagnostic assay, to detect antigens and antibodies by changes in enzyme activity.
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/note= "this position is linked to the OMPC of the
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                                                                                                                                                                                                   80.5%; Score 62; DB 2; Length 13; 100.0%; Pred. No. 0.015; ive 0; Mismatches 0; Indels
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                                       Claim 8; Page 88; 133pp; English
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90US-00555966.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Leanza WJ,
                                                                                                                                                                                                                                                                                                                                                                             AAR31254;
                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                         RESULT 191
                                                                                                                                                                                                                              Matches
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us-08-869-386-1.rag

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or protein complex having a first set of covalent linkages to low molecular weight moieties which have an anionic or polyanionic character at physiclogical ph, and a second set of covalent linkages to peptides comprising HIV principal neutralizing determinants (PND's) or immunologically equivalent peptides. Preferably at least one set of the immunologically equivalent peptides. Preferably at least one set of the covalent linkages is comprised of maleimide derivatives; the covalent linkages is comprised of maleimide derivatives; the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria protein is the outer membrane protein complex (OMPC) of Neisseria bonded cyclic structure, an amide-bonded cyclic structure or a thioether-conded cyclic structure. The present sequence (cPND3) is an example of a PND peptide component used in the co-conjugate. The co-conjugate is PND peptide component used in the co-conjugate. The co-conjugate is useful for inducing anti-peptide immune response in mammals, for inducing HIV-neutralizing antibodies in mammals, for Eormulating vaccines to gardilicted with HIV infection or disease
                                                                                                                                                                                                                                                                                                                                                                  ö
The invention relates to a co-conjugate comprising an immunogenic protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note= "forms cyclic thio ether linkage with Cys(15)"
                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "forms cyclic thio ether linkage with Cys(2)"
                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PND; thio-ether cyclic linkage; HIV; AIDS; assay; vaccine.
                                                                                                                                                                                                                                                                                                                                  80.5%; Score 62; DB 2; Length 15;
100.0%; Pred. No. 0.017;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cyclic HIV principal neutralising determinant peptide.
                                                                                                                                                                                                                                                                                                                                            100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR20214 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90US-00555227.
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                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   1 RIQRGPGRAFVT 12
                                                                                                                                                                                                                                                                                                                                                                                                                     (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hannah J, Tolman RL;
                                                                                                                                                                                                                                                                                                                                                 Local Similarity
nes 12; Conser
                                                                                                                                                                                                                                                                                                      Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Modified-site
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Matches
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are bridged by o-xylylene, giving a ring system having a stable thiocether bond. The peptide is a preferred example of a more generic molecule given in Claim 1, comprising the "GPGR- core sequence as part of a peptide ring system completed by a thio-ether bond formed by linking the omega-mercapto groups of two Cys or homologous residues via xylylene ring is more stable than that formed using a disulphide bond. The peptides are useful as analytical tools, as reagents in ELISA assays, or as reagents for maining covalent conjugate immunogens. The conjugates are useful for raising mammalian anti-peptide, anti-HIV or HIV-neutralising immune responses and can be used as vaccines and therapeutics for AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 via o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "forms cyclic thioether linkage with Cys(2) via o-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "bonded via N-terminal to polysaccharide- protein complex"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a conjugate of an HIV principal neutralizing
                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus; principal neutralizing determinant; outer membrane protein complex; OMPC; Neisseria; AIDS; cyclic; cPND3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV peptide-polysaccharide-protein conjugates - used in vaccines or produce antibodies to prevent or treat HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "forms cyclic thioether linkage with Cys(15)
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                                                                                                                                                                                               Length 15;
                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV-PND-polysaccharide-protein conjugate vaccine
                                                                                                                                                                                              Score 62; DB 2;
Pred. No. 0.017;
                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                    AAR26689 standard; peptide; 15 AA.
                                                                                                                                                                                     80.5%; Sur
100.0%; Pre
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91US-00715275.
91US-00715277.
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                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                               3 RIQRGPGRAFVT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Marburg S, Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1992-034437/05
                                                                                                                                                                                                             Local Similarity
les 12; Conserv
                                                                                                                                                                       Sequence 15 AA;
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Modified-site
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19-7UL-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                               AAR26689;
                                                                                                                                                                                                Query Match
                                                                                                                                             and ARC
                                                                                                                                                                                                                                                                                                                          RESULT 193
                                                                                                                                                                                                                            Matches
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New cyclic HIV principal neutralising determinant peptide(s) - with of stable thio-ether bonds used as reagents in enzyme-linked immunosorbent assays or conjugates in vaccines for HIV and AIDS.

WPI; 1992-026504/04.

The omega-mercapto groups of the Cys residues at the 2 and 15 positions

Claim 4; Page 13; 16pp; English

Gaps

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Indels

Pred. No. 0.017;

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12; Conservative 1 RIQRGPGRAFVT 12 3 RIORGPGRAFVT 14

Local Similarity

Best Loc Matches

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100.08;

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determinant (PND), or an immunologically equivalent peptide (PEP), covalently coupled to an immunogenic protein complex through an anionic polysaccharide linker. Pref. the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b and the PND peptide has a linear structure, a disulphide-bonded cyclic structure. The present sequence (CPND3) is an example of a PND peptide component. The mercapto groups of Cys(2) and Cys(15) are bridged by o-xylylene. The marking vaccines to prevent contraction of HIV infection or disease. The antibodies can be used for inducing HIV-neutralising antibodies or for antibodies can be used for protecting against infection by HIV, or for protecting against proliferation of HIV post-infection by HIV, treating AIDS, or in diagnostic assays
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                                                                                                                                                                                                       80.5%; Score 62; DB 2; Length 15; 100.0%; Pred. No. 0.017; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Alkaline phosphatase; AP; amino acid moiety; AAM.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Alkaline phosphatase 91-93 replacement.
                                                                                                                                                                                                                                                                                                                                                AARS8606 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94WO-US002539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93US-00031165
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(first entry)
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Matches 12; Conservative
                                                                                                                                                                                                                                                                                  3 RIORGPGRAFVT 14
                                                                                                                                                                                                                                                           1 RIORGPGRAFVT 12
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                                                                                                                                                                                 Sequence 15 AA;
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01-MAY-1995
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Gaps

AARS8606 is an amino acid molety (AAM), when it replaces the amino acid residues 91-93 of alkaline phosphatase (AP) (AARS8608) it enables the enzymatic activity of AP to be modulated upon the binding of a molecule to the inserted sequence. The modified AP can now be used as a diagnostic assay, where the presence of specific antigens and antibodies can be inferred by changes in the enzymes activity. (Updated on 25-MAR-2003 to Genetically engineered hybrid enzymes and ligand conjugates - useful for diagnostic assay, to detect antigens and antibodies by changes in enzyme Claim 8; Page 88; 133pp; English, activity.

Mandecki

Krafft GA,

Bridon DP, Jaffe KD, Yue VT;

Brate EM, Brennan CA, March SC, Russell JC, WPI; 1994-303041/37.

Sequence 15 AA;

correct PN field.)

Query Match

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DB 2; Length 15;
80.5%; Score 62;
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epitope recognized by NM-01, the antibody was screened by ELISA for epitope recognized by NM-01, the antibody was screened by ELISA for eactivity with overlapping peptides corresp. to the amino acid sequence of the V3 loop region of HIV-1 gpl20 (AAR33332, AAR3333, AAR3333).

While there was no detectable reactivity over background of MAb-01 with the peptides corresp. to AAB 302-316 or 322-336 of the V3 loop, binding of the antibody to the peptide representing AAB 3122-326 was apparent.

The extent of this reactivity with other HIV-1 isolates was screened with peptides corresp. to the V3 loop region of HIV-1 isolates IIIB, RF, CDC4, NV/5, Z6, Z2 and ELI (AAR33335-R33342). These results indicate that monoclonal antibody NM-01 recognizes an epitope of the V3 loop of gpl20 of multiple HIV-1 isolates having the amino acid sequence AAR33334.

CM woll were cloned by PCR and sequenced. Nucleotides 1-21 and 334-363 of AAQ37471 corresp. to the PCR primers used to amplify NM-01 light chain sequences and nucleotides 1-27 and 385-402 of AAQ37471 corresp. to the CCC MAR-2003 to correct PN field.)
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                                                                                                                                  equence of peptide which corresp. to the V3 loop region of gp120 of HIV-isolate IIIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monoclonal antibodies against HIV-1 gp120 and gp160 proteins - for treating and preventing HIV-1 infection.
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Pred. No. 0.031;
1; Mismatches 1; Indels
                                                                                                                                                                                         Monoclonal antibody; NM-01; HIV-1; gp120; gp160.
                                                                                                                                                                                                                                                                                                                                                                                   (NISP ) NISSIN SHOKUHIN KAISHA LTD.
                                 AAR33336 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 20; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                       91US-00748562.
                                                                                                                                                                                                                                                                                                                         92WO-US007111.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 IQRGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12; Conservative
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Matches 12; Conserv
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06-JUL-1993
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                                                                                                                                                                                                                           Synthetic.
                                                               AAR33336;
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RESULT 195
                  AAR33336
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The invention relates to a novel monoclonal antibody designated NM-01. The antibody was raised by immunising 2-month old Balb/c mice with live thu. Splenocytes from the mice were fused to P3-K63-Ag8-U1 cells (ATCC CRL1597). Hybridomas were screened using membranes from non-infected and MN-infected H9 cells, by reacting with hybridoma culture supernatants. This screening was followed by immunofluorescence and radioimmunoassays. The screening isolated the hybridoma HB 10726 which secrees the antibody NM-01. The peptides AAW09263-72 are derived from other HIV strains and were used to determine which other HIV-1 isolates antibody NM-01 reacted with. This peptide is from HIV-1 strain IIIB. The antibody is used for the diagnosis of HIV-1 in a fluid e.g. blood, and can be used to treat or prevent an HIV-1 infection. (Updated on 25-MAR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New monoclonal antibodies to HIV-1 - used for the prevention, treatment or diagnosis of {\rm HIV}{-1} infection.
                                                                                                                                                                             Human immunodeficiency virus type-1; HIV-1; gpl20; gpl60; epitope; monoclonal antibody; infection; heavy chain; light chain; hybridoma; complementarity determining region; CDR; V3 loop.
                                                                                                                                    HIV-1 strain IIIB gp120 V3 loop peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NISP ) NISSIN SHOKUHIN KAISHA LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          91US-00748562.
92WO-US007111.
93US-00039457.
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                                                                                         (first entry)
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Best Local Similarity 85.7
Matches 12; Conservative
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                                                                     (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                    24-AUG-1993;
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22-APR-1993;
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03-MAR-1997
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                        AAW09264;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 198
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셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GPCR is a portion of HIV-1 gpl20 or gpl60 protein. Monoclonal antibodies (MAbs) that react with this and which have the capacity to neutralise the infection of H9 cells in culture by live HIV-1 strains MN and IIIB are claimed. Specifically illustrating the invention are the murine MAb (designated NM-01) produced by hybridoma cell line HB 10726 which is deposited under ATCC No. HB 10726, and the humanised versions of Ab NM-01. To identify the specific epitope of gpl20 recognised by NM-01, the Ab was screened for reactivity with three overlapping peptides corresp. to the V3 loop region of gpl20 (AAR48600-02). While three was no detectable reactivity over background of MAb NM-01 with the peptides corresp. to AAs 102-316 of the V3 loop, binding of the Ab to the peptide corresp. to binds to the V3 loop region of HIV-INW gpl20 prompted further studies on the extent of this reactivity with other HIV-I isolates. The Ab was creened by ELISA for reactivity with other HIV-I isolates. The Ab was seriened by ELISA for reactivity with peptides corresp. to the AB sequences of the peptides are given in AAR48603-10. NM-01 reacted with especial or HIV-I isolates IIIB, RF, and CDC4 isolates. It showed a lesser affinity for the NY/5 peptide. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                      Human immunodeficiency virus; HIV-1; AIDS; glycoprotein; V3 loop; gpl20;
epitope; isolate IIIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New humanised antibody specific for epitope on HIV-1 gp 120 - able to neutralise infection of HG cells, also nucleic acid encoding it, useful for passive immunisation to treat or prevent HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                               Sequence of portion of gp120 V3 loop peptide from HIV-1 isolate IIIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 60; DB 2; Length 14;
Pred. No. 0.031;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example; Table 4, Page 18; 91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NISP ) NISSIN SHOKUHIN KAISHA LTD.
                                                                AAR48604 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93WO-US007967.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93US-00039457.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             77.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92WO-US007111
                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 IORGPGRAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IRIGPGRAFVTICK 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 85.7
hes 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1994-083117/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                            WO9404574-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-AUG-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-AUG-1992;
                                                                                                                                                         25-MAR-2003
03-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                         03-MAR-1994
                                                                                                              AAR48604;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ohno T;
                     RESULT 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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AAR4 864

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                      Gape
                                                                                                                                                                                                         HIV-1 gp120 V3 loop domain containing U1 snRNP 70K consensus epitope.
                      ö
77.9%; Score 60; DB 2; Length 14; 85.7%; Pred. No. 0.031;
                      1; Indels
                                                                                                                                                                                                                                epitope; autoantibody; immunoinfective cluster virus;
                      1; Mismatches
                                                                                                                                                                                     (first entry)
                                                                                                                                                                                      03-MAY-1995
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AAW09264 standard; peptide; 14 AA.

RESULT 197 AAW09264 ID AAW092

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New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency

(AMNA-) AMERICAN NAT RED CROSS

Scott D, Zambidis E;

WPI; 1998-506315/43.

97US-0040581P. 98WO-US002766

13-FEB-1998; 13-FEB-1997; Claim 10; Page 119; 154pp; English

virus infection.

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Previous immunological analyses of the V3 loop of HIV-1 (AAR62159) have localised the main neutralising domains. The target of more than 80% of neutralising antibodies in HIV-1 infected sera from AIDS patients has now been found to overly the consensus binding sequence and domain A epitopes of the U1 snRNP 70K protein. In AIDS, antibody titres are too low to immunodominant targets of autoantibodies sequences in 70K are immunodominant targets of autoantibodies in the systemic rheumatoid disorder of mixed connective tissue disease. The titrers of such disorder of mixed connective tissue disease. The titrers of such cross-react with HIV-1 epitopes and are useful for treating HIV infection. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                             Methods for treating immunoinfective cluster virus infections - utilise antibodies or fragments characteristic of auto antibodies produced by patients with rheumatic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
             human immunodeficiency virus, HIV-1; systemic lupus erythematosus, mixed connective tissue disease; scleroderma; glycoprotein 120; Ul snall nuclear ribonucleoprotein; Ul snRNP 70K protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                75.3%; Score 58; DB 2; Length 11; 100.0%; Pred. No. 0.051; 1.ve 0; Mismatches 0; Indels
nuclear protein antigen; systemic rheumatic disorder;
                                                                                                                                                                                                                                                                                  Ehresmann G;
                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure, Page 62, 106pp, English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW76852 standard; peptide; 11 AA.
                                                                                                                                                                                                                                               (UYSC-) UNIV SOUTHERN CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.33,
100.0%; Pre-
                                                                              Human immunodeficiency virus 1.
                                                                                                                                                                                  94WO-US002631.
                                                                                                                                                                                                                 93US-00029850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-JAN-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 RGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                  Douvas A, Takehana Y,
                                                                                                                                                                                                                                                                                                                WPI; 1994-302689/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11 AA;
                                                                                                                                                                              10-MAR-1994;
                                                                                                              WO9420141-A1
                                                                                                                                                                                                                11-MAR-1993;
                                                                                                                                               15-SEP-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9836087-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-AUG-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW76852;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8
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This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) procein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immunoglobulin heavy virus (HHV) gp120 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gp120 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gp120, so prevents or inhibits 'bystander' apoptosis of uninfected T cells that are bound to gp120 protein, maximising induction of protective antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polyfunctional base sequence; microgene; industrial; cell culture; artificial matrix protein; transgenic animal; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     75.3%; Score 58; DB 2; Length 11; 100.0%; Pred. No. 0.051; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV gp120 related peptide SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NISC-) JAPAN SCI & TECHNOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABB05777 standard; peptide; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LS-JUN-2001; 2001WO-JP005116.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-JUN-2000; 2000JP-00180997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 RGPGRAFVTIG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200196558-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-AUG-2003
07-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABB05777;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shiba K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Gaps

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Gaps

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Length 13; 1; Indels

DB 5;

75.3%; Score 58; DB 5; 91.7%; Pred. No. 0.059; 0; Mismatches

11; Conservative

Matches

Query Match Best Local Similarity

Sequence 13 AA;

us-08-869-386-1.rag

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Unidentified
                                                 25-APR-2002
                     Query Match
                                   AA015659
                                                         Shiba K,
   culture.
                       Matches
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                                                                                                                                                                                             The present invention describes a polyfunctional base sequence (NI) having two or more functions in different reading frames. Also described are: (1) a method for producing NI and artificial gene expression vectors comprising NI; (2) transgenic non-human animals comprising NI; and (3) tracements and disgnostic reagents containing an artificial protein, artificial tissues or high molecular weight artificial proteins. In useful for creating industrially useful artificial matrix proteins for call culture. The present sequence represents a peptide which is used in example from the present invention. (Updated on 29-AUG-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Strong immune response induction; high-order protein structure formation;
Polyfunctional base sequence having two or more functions in different reading frames, useful for producing artificial matrix proteins for cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Artificial protein capable of inducing a strong immune response to a peptide group for assisting antibody production in vivo to viruses and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 5; Length 13;
0.059;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Strong immune response induction-related peptide 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 58;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NISC-) JAPAN SCI & TECHNOLOGY CORP.
                                                                                                                                      Example 1; Page 47; 61pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 46; 77pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAO15659 standard; peptide; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      75.3%;
91.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIQRGPGRAFVT 12
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RIQRGPGRTFVT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200233074-A1.
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The invention comprises an artificial protein which induces a strong immune response to a peptide group (the protein contains all or part of the peptide group). The artificial protein assists the formation of high-order protein structure and/or assists the antigen presentation of immunocompetent cells. The artificial protein of the invention is useful for inducing a strong immune response and the preparation of effective antibodies to specific antigens, especially HIV. The present amino acid sequence represents a peptide that was used in the invention

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racing sources of infectious diseases. The method comprises preparing a short (9-50 amino acid) peptide containing at least one non-conserved short (9-50 amino acid) peptide containing at least one non-conserved intradermally into a test subject in a delayed-type hypersensitivity (DTH) assay and observing the injection site at intervals for induration. In this example CD4+T-lymphocyte epitopes to the HIV-1 V3 loop were mapped by existing in vitro technique for two existing HIV infected individuals and used to design peptides AAW24217-20. The method allows the T-Tymphocyte epitopes of a large antigen to be determined in vivo in humans. The method is useful in medicine e.g. in diagnosis, monitoring and treatment design for infectious disease exposure, active autoimmune disease, allergic diseases and malignancy. It is sepecially useful for tracing infectious diseases e.g HIV, particularly when a sequence is present only in certain strains of an organism, and developing suitable vaccines. Vaccinated individuals can also be tested to verify protection against a particular strain. The method allows in vivo mapping of Tenymphocyte epitopes, not previously possible. The method is simpler, more rapid and more sensitive. It can also be applied in a variety of environments e.g. undeveloped regions since specialist equipment is not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide fragments containing antigen epitope(s) used to trace diseases used in a delayed-type hypersensitivity assay, for in vivo mapping of human T-lymphocyte epitope(s) e.g. for diagnosis, vaccine development
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Mapped CD4+T-lymphocyte epitope of patient 2"
                                                                                                                                                                                                                                                         CD4+ T-lymphocyte epitope to HIV-1 V3 loop derived peptide V3-LAI-B.
                                                                                                                                                                                                                                                                                              T-lymphocyte epitope; diagnosis; antigen; infectious disease; delayed-type hypersensitivity assay; vaccine development.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (USSA ) US DEPT ARMY GOVERNMENT US ARMY MEDICAL
                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure, Page 6; 14pp; English.
                                                                                                                                          AAW24218 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97WO-US001084.
                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96US-0010679P.
                                                                                                                                                                                                                      (first entry)
1 RIORGPGRAFVT 12
                                         RIORGPGRIFVT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sitz KV, Brix DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9727462-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-JAN-1996;
                                                                                                                                                                                                                      17-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-JUL-1997
                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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                                                                                                     RESULT 202
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cytotoxic T lymphocyte; epitope; antigen; pathogenic; nef; gag; pol; env gpl20; gp41; HIV; cell-mediated immunity; human immunodeficiency virus; class II restricted.

Human immunodeficiency virus.

W09428871-A1

94WO-US006394. 93US-00072718.

07-JUN-1994;

22-DEC-1994.

(ENDO-) ENDOCON INC.

Leonard RJ;

07-JUN-1993;

WPI; 1995-036067/05.

Cytotoxic T lymphocyte epitope 56 derived from env gp120 protein.

(revised)
(first entry)

25-MAR-2003 23-AUG-1995

AAR68799 standard; peptide; 11 AA.

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AAR68799
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nine peptides from the V3 regions of HIV-1 isolates IIIb, RF and MN were incorporated into terravalent multiple antigen peptide systems (MAPS) (see AAR32406-14). Parallel groups of three peptides with chain lengths spanning from 11-24 residues were synthesised in MAPS formate for each isolate. ELIS assays demonstrated that antisera titers in mice were immunisation - the length of the IIIb peptide used for the closely related to the length of the IIIb peptide used for the substantial antibody prodn. in mice against the other two series of substantial antibody prodn. in mice against the other two series of speptides, RP (B4-B6), and MN (B7-B9), except for a low reactivity in the gp. immunised with B8 (MN isolate). Specificity tests of the B cell response demonstrated that the T cell epitope (AAR22415) also serves as a B cell epitope. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence of peptide B3 which comprises AAs 315-325 from the V3 region of HIV-1 isolate IIIb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New multiple antigen peptide(s) as HIV vaccines - include a dendritic core covalently bonded to peptide including the sequence IGPGR.
                                                                                                                 Gaps
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                                                            75.3%; Score 58; DB 2; Length 19; 100.0%; Pred. No. 0.083; ive 0; Mismatches 0; Indels
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100.0%; Pred. No. 0.072;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV-1; vaccine; dendritic core; ss.
                                                                                                                                                                                                                                                                                                                            AAR32408 standard; peptide; 11 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92WO-US006688
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(first entry)
                                          Query Match
Best Local Similarity 100.
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(UYRQ ) UNIV ROCKEFELLER.
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Best Local Similarity 100.
Matches 11; Conservative
                                                                                                                                                                                    5 GPGRAFVTIGK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tam JP, Profy AT;
               Sequence 19 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 11 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-MAR-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003
04-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                              AAR32408;
                                                                                                                                                                                                                                                                               RESULT 203
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AAR68744-805 are cytotoxic T lymphocyte (CTL) class I and II restricted epitopes derived from human immunodeficiency virus proteins. AAR68799 corresponds to amino acid residues 310-316 of the env gpl20 protein. These antigens are examples of peptides that can be used with an immunogenic implant. The implant is associated with an antigen associated with a pathogen and used to form a discrete, localised chronic inflammation site which acts as a local 'factory' for prodn. of CTL's pathogen-specific CTL's an eradicate or prevent development of inflection, and can also be used to tract or arrest the development of cancers associated with infection. (Updated on 25-MAR-2003 to correct PN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Glutamic acid; glutamine; vaccine; major histocompatibility complex; MHC; immunomodulator; antiallergic; endocrine; neuroprotectant; virucidal; bactericidal; antiparasitic; fungicidal; cytostatic; medicine;
Implant for sustained release of pathogen-associated antigen - forming chronic inflammatory site producing cytotoxic T-lymphocytes lysing infected cells, esp. for treating AIDS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vaccine related MHC ligand peptide SEQ ID NO:533.
                                                                                                                Disclosure, Page 12; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAM99430 standard; peptide; 11 AA.
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Gaps

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RESULT 204

Length 11;

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The present invention describes a pharmaceutical compound (I) that contains an N-terminal glutamic acid (Glu) or glutamine (Gln) residue in the form of an addition salt with a strong, physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition containing at least one (I); (b) a vaccine containing at least one (I) where this is a major histocompatibility complex (MFO) ligand (Ia); (c) a method for in vitro diagnosis of diseases associated with the presence of (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process (Ia); a kit for method (c) that includes a (Ia); and (e) a process (II); an euseful, in human or veterinary medicine, repreparing (I). (I) has immunomedulator, endocrine; e.g. in pharmaceutical compositions (for treating immune disorders, e.g. immune deficiency, autoimmune states, hypersensitivity, allergy, graft rejection, infection, otherwise (I) is a MHC ligand (Ia), in vaccines for treatment or prevention of: (I) viral, bacterial, parasitic or fungal infections; or (ii) of cancers. A particular application is in antimelanoma vaccines. (I) are also useful for in vitro diagnosis of diseases associated with interactions between MHC and (I), e.g. melanoma and human immunodeficiency virus infection. AMM98899 to AAM99592 represent invention control of the control of menucodeficiency virus infection. AMM98899 to AAM99592 represent invention
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pharmaceutical; immune disorder; immune deficiency; autoimmune; hypersenakitvity; allergy; graff rejection; infection; hormonal disorder; central nervous system disease; cancer; melanoma; anti-melanoma vaccine; human immunodeficiency virus.
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vif; tat; cytotoxic T lymphocyte; CTL; immune response; epitope; antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Stabilized pharmaceutical containing N-terminal glutamic acid or glutamine, useful e.g. in anti-melanoma vaccines, is an addition salt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Pred. No. 0.072;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                     Goetsch L;
                                                                                                                                                                                                                                                                                                                                                     Klinguer-Hamour C, Corvaia N, Beck A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV B27 super motif env peptide #127.
                                                                                                                                                                                                                                                                                                            (FABR ) FABRE MEDICAMENT SA PIERRE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              74.0%; SCOL.
100.0%; Pre
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 122; 149pp; French.
                                                                                                                                                                                                                             22-MAR-2001; 2001WO-FR000872.
                                                                                                                                                                                                                                                                   23-MAR-2000; 2000FR-00003711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
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nes 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             with strong acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11 AA;
                                                                                                                                         WO200170772-A2
                                                                                                    Mus musculus.
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15-JUL-2002
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The present invention to describe a composition, or compilating an amino acid sequence selected from 51 defined amino acid sequences (ABL25347 to ABP25397). (1) has virtucide activity and can be used in vaccines. (1) may be used for immunishing subjects against HIV-1 infections. The use of group-based vaccines has several advantages over traditional vaccines. Compositions. There is evidence that the immune response to whole compositions. There is evidence that the immune response to whole antigens is directed largely toward variable regions of the antigen, and proup-based vaccine may be selected from conserved regions of viral or tumour-associated antigens, which therefore reduces the likelihood of the uniquence of the antigens, which therefore reduces the likelihood of the uniquence of antigens, which therefore reduces the likelihood of the natigens can be avoided with the use of group based vaccines. An additional advantage of an group-based vaccine approach is the ability to combine selected groups (CTL and HTL), and further, to modify the composition of the groups, achieving, for example, enhanced is municated. Accordingly, the immune response can be modulated, as immunogenicity. Accordingly, the immune response can be modulated, as immunogenicity. Accordingly, the immune response can be modulated as in not possible with traditional approaches. ABP11501 to ABP25412 represent peptide sequences used in the examplification of the present invention. (Updated on 11-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a composition (I) comprising a prepared
                                                                                                                                                                                                                                                                                                                                                                         Vaccine compositions comprising human immunodeficiency virus-1 (HIV-1) peptide groups, useful for vaccinating against HIV-1.
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                                                                                                                                                                                                                                                                            Cheanut
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              74.0%; Score 57; DB 4; Length 11; 100.0%; Pred. No. 0.072; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                            Livingston BD,
vaccine, HIV infection; immunisation; virucide
                                                                                                                                                                                                                                                                            Southwood S, Livi
Kubo RT, Grey HM;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 32; Page 219; 448pp; English.
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                                       Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protease inhibitory peptide #1
                                                                                                                                                      05-OCT-2000; 2000WO-US027766.
                                                                                                                                                                                                05-OCT-1999; 99US-00412863.
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                                                                                                                                                                                                                                       EPIM-) EPIMMUNE INC
                                                                                                                                                                                                                                                                            Sette A, Sidney J,
Baker DM, Celis E,
                                                                                                                                                                                                                                                                                                                                        WPI; 2001-354887/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                            WO200124810-A1
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                                                                                                                     12-APR-2001.
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Matches
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Synthetic

localised the main neutralising domains. The target of more than 80% of neutralising antibodies in HIV-1 infected sera from AIDS patients (AAR62164) has now been found to overly the consensus binding sequence and domain A epitopes of the UI snRNP 70K protein. In AIDS, antibody titres are too low to arrest the disease; however, the homologous sequences in 70K are immunodominant targets of autoantibodies in the systematic rheumatoid disorder of mixed connective tissue disease. The titers of such autoantibodies exceed 10 power 7. The anti-snRNP autoantibodies will cross-react with HIV-1 epitopes and are useful for treating HIV infection. (Updated on 25-MAR-2003 to correct PN field.)

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Gaps

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74.0%; Score 57; DB 2; Length 15; 100.0%; Pred. No. 0.095; ive 0; Mismatches 0; Indels

11; Conservative 1 RIORGPGRAFV 11 5 RIORGPGRAFV 15

Best Local Similarity

Matches

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Query Match

Sequence 15 AA;

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                                                                                                                                                                                 A protease inhibitor which exhibits an improved inhibitory effect comprises this peptide or its salt. The peptide must include at least the tetrapeptide GPGR and does not contain the sequence NNTRKSIRIRIQRGPGRA.
                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      epitope; autoantibody; immunoinfective cluster virus;
nuclear protein antigen; systemic rheumatic disorder;
human immunodeficiency virus; HIV-1; systemic lupus erythematosus;
mixed connective tissue disease; scleroderma; glycoprotein 120.
                                                                                                                            New peptide protease inhibitor - used as hapten for prodn. of immunoglobulin(s) and as antiinflammatory antitumour agent etc.
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                                                                                                                                                                                                                                                           74.0%; Score 57; DB 2; Length 12; 100.0%; Pred. No. 0.078; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-1 gp120 V3 loop neutralising domain.
                                                                                                                                                                                                                                                                                                                                                                              AAR62164 standard; peptide; 15 AA.
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                                                                                                                                                            Claim 1; Page 1; 9pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus 1.
                                           89JP-00135678.
                                                               89JP-00135678
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                                                                                    (NITL ) NITTO DENKO CORP.
                                                                                                                                                                                                                                                                                 11; Conservative
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                                                                                                         WPI; 1991-048837/07.
                                                                                                                                                                                                                 See also AAR10593-4
                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                       Sequence 12 AA;
JP03002195-A
                                        29-MAY-1989;
                                                              29-MAY-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-MAR-1994;
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                     08-JAN-1991,
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03-MAY-1995
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New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
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                                                                                                                                                                        B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                         Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 10; Page 119; 154pp; English.
                                  AAW76846 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AMNA-) AMERICAN NAT RED CROSS.
                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-US002766.
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                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-506315/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virus infection
                                                                                                                                                                                                                                                                                                                                                 WO9836087-A1.
                                                                                                           25-JAN-1999
                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                   20-AUG-1998.
                                                                      AAW76846;
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RESULT 209
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                   AAW76846
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Methods for treating immunoinfective cluster virus infections - utilise antibodies or fragments characteristic of auto antibodies produced by patients with rheumatic disorders.

Douvas A, Takehana Y, Ehresmann G;

WPI; 1994-302689/37.

Previous immunological analyses of the V3 loop of HIV-1 (AAR62159) have

Disclosure; Page 61; 106pp; English.

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AAR25139 standard; protein; 17 AA.
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30-AUG-1995
                                                                                                 25-MAR-2003
05-JAN-1993
                                                                                                                                                                                                                                           5-JUN-1992
                                                                                                                                                                                          Synthetic.
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                                                                          AAR25139;
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                         RESULT 211
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Induction of tolerance suppresses production of antibodies against gp120, so prevents or inhibits 'bystander' apoptosis of uninfected T cells that are bound to gp120 protein, maximising induction of protective antiviral T cell immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            In vitro antigen specific activation of antibody producing cells comprises culturing a population of isolated, non-adherent mononuclear immune cells for a time and under conditions sufficient to induce differentiation of the cell.
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                                                                                                                                                                                                                                                                                                          Peptide epitope from HIV-1 gp120 protein corresponding to aa 304-318
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                                                                                      74.0%; Score 57; DB 2; Length 15;
100.0%; Pred. No. 0.095;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                  antigen specific activation; antibody producing cell; non-adherent mononuclear immune cell; T helper cell; lysosome-containing cell; differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 29; SEQ ID NO 4; 88pp; English.
                                                                                                                                                                                                                              ADP76013 standard; peptide; 15 AA.
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                                                                                                                  11; Conservative
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                                                              Sequence 15 AA;
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                                                                                       Query Match
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RIORGPGRAFV 11

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RNA mol. derived from alphavirus RNA genome - chimeric alphavirus antigen and vaccine for immunisation against viral infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T-cell; epitope; HIV-1; core protein; p24E; B-cell; antigen; gp160; gag; pol; vaccine; multimeric peptide; AIDS; 3D organisation.
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Pred. No. 0.11;
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                                                                                                         Semliki forest virus; SFV; E2 protien; vaccine
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                                                                 SFV-HIV epitope
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                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents the a hybrid B-cell epitope derived from the HIV epitope. 193 (BRU) and (MN) loops. This peptide can be ligated to a T cell epitope, eg. p24E (see also AAR6851) to form a chimeric peptide. Chimeric peptides such as this, may then be used in the production of multimeric peptides such as this, may then be used in the production of multimeric peptides sequences may also be used in the production of multimeric peptides in which the peptides are C-terminally modified by the carry an additional copy of the peptide modified on its epsilon amino acid multimeric peptides may be used for the treatment of AIDS by acting to displace the binding of HIV virus to human or animal cells or by disturbing the 3D organisation of the virus. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                          Novel tandem synthetic HIV-1 peptide(s) - comprising T-cell epitope of gag protein linked to B-cell epitope of V3 loop protein of an HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV; human immunodeficiency virus; gag; T-cell; B-cell; epitope; env; V3 loop; vaccine; determinant; chimaeric.
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/note= "Residues 315-325 of the V3 (MN) loop"
                  /note= "Residues 307-316 of the V3(BRU) loop"
                                                                                                                                                                                                                                                                                                                                                                                                 2; Length 21;
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Pred. No. 0.13;
); Mismatches
 Location/Qualifiers
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                                                                                                                                                                                                                                 Disclosure; Page 12; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW25815 standard; peptide; 21 AA.
                                                                                              94WO-CA000317
                                                                                                                 93US-00073378
                                                                                                                                                      Chong P, Klein MH
                                                                                                                                                                                                                                                                                                                                                                                                74.0%;
91.7%;
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(first entry)
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                                                        WO9429339-A1.
                                                                                              08-JUN-1994;
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20-OCT-1997
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Key
Peptide
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amino acid sequence comprising an HIV gag protein T-cell splitope linked epitope of the V3 loop of an HIV gag protein T-cell splitope linked epitope of the V3 loop of an HIV env protein, which can be used to generate vaccines against HIV-I. The T-cell epitope sequence is pref. Selected from the T-helper determinant core peptides P24B, P24B, P24L, P24M and P24H while the B-cell epitopes are derived from HIV strains including CTLB-56, V3NM, CTLB-59, CTLB-55, SF2, LAI, IIIB, RF, Z6, Z054, LIV14 and BXOB. The peptides are chimmeric and can be linked to a branched Lys backbone. This sequence represents a hybrid HIV-1 envy protein V3 loop B-cell epitope comprising amino acids 307-316 of the V3 loop from the BNU strain (AAM25816) linked at its C-terminus to amino acids 315-325 of the MN strain V3 loop (AAW25817). The VP peptide is linked to the C-terminus of the p24B peptide to generate the chimmeric peptide VP-T-B (AAW25814). (Updated on 25-MAR-2003 to correct PF field.)
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                                                                                                                                                                                                                                                                                                                             The invention relates to new synthetic peptides comprising at least one
                                                                                                                                                                             Tandem synthetic HIV peptide(s) useful as immunogens - comprising gag protein T-cell epitope linked to env protein B-cell epitope.
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Pred. No. 0.13;
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                                                              Chong P;
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91.7%;
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(CONN-) CONNAUGHT LAB LTD.
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09-JUN-1994;
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                                                           Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Synthetic human immunodeficiency virus-1 peptide(s) - containing T-cell epitope and B-cell epitope(s) are candidate vaccines against  $\rm HIV-1.$ 

Chong P, Klein MH, Sia CDY;

WPI; 1998-556461/47

(CONN-) CONNAUGHT LAB LTD.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a synthetic peptide comprising an amino acid sequence containing a T-cell epitope of an HIV gag protein linked at its C terminus to an amino acid sequence containing a B-cell epitope of an HIV gp41 protein and containing the amino acid sequence: XILKDWX2; where XI = E, A, G or Q, and X2 = A or T, or an amino acid sequence capable of eliciting an HIV-specific antiserum and recognizing the sequence XILKDWX2. The synthetic peptide is useful in vaccines against HIV infection and in diagnostic applications. AAW99892 to AAW99906, and AAW99999 to AAW99989 represent synthetic peptides from the present
                              The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-cell epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitopes are generally designed based on the p24 core protein and the B-cell epitopes from the V3 loop of the gpl20 protein from various HIV-1 strains. This peptide corresponds to a hybrid B-cell epitope which is a chimera of the B-cell epitopes V3 (BRU) and V3 (MN) (AAW67332-W67333), respectively. The peptide is used to generate the chimeric T- and B-cell epitope VP-T-B (AAW67330)
                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic chimeric HIV polypeptides – comprising gag protein T-cell epitope linked to gp41 B-cell epitope.
                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV-1; human immunodeficiency virus; vaccine; T-cell epitope;
gag protein; B-cell epitope; gp41 protein; chimeric; infection.
                                                                                                                                                                                                                                         74.0%; Score 57; DB 2; Length 21; 91.7%; Pred. No. 0.13; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV-1 vaccine synthetic peptide SEQ ID NO:16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Col 33-34; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW99939 standard; peptide; 21 AA.
 Disclosure; Col 7; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95US-00462507.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93US-00073378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94US-00257528
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                                                                                                                                                                                                                                                             Local Similarity 91.7
nes 11; Conservative
                                                                                                                                                                                                                                                                                                                1 RIQRGPGRAFVT 12
                                                                                                                                                                                                                                                                                                                                                   7 RIORGPGRAFYT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-189590/16
                                                                                                                                                                                                            Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US5876731-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention
                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 215
                                                                                                                                                                                                                                                                                Matches
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DB 2; Length 21;

74.0%; Score 57;

Sequence 21 AA;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents a fragment of a HIV1 protein, and can be used in the immunogenic composition of the invention. The composition comprises a synthetic fusion polypeptide which includes a sequence encoding 1 or more T cell epitopes and a sequence encoding 1 or more B carrier. Both the T cell and B cell epitopes are derived from HIV proteins. The compositions are useful as vaccines against HIV infection. The composition induces HIV-1-specific polyclonal antibodies that are opponising and antiviral. The specific polyclonal antibodies that are induce a response against different viral isolates and in subjects who recognise different T cell epitopes. (Updated on 17-OCT-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency virus, useful antigens in producing vaccines.
                                                                                                                                                                                                                                                                         composition; T cell epitope; B cell epitope;
              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                Indels
              1;
 Pred. No. 0.13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                             AAY39690 standard; peptide; 21 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR66418 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Col 7; 43pp; English.
                                                                                                                                                                                                                                              HIV1 gag protein epitope VP-p24.
              ö
                                                                                                                                                                                                                                                                                      infection; antibody; antiviral
                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sia CDY;
                                                                                                                                                                                                                                                                                                                                                                                                          95US-00467881.
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94US-00257528.
 l Similarity 91.7%;
11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                      (revised)
(first entry)
                                                                                                                                                                                                                                                                            vaccine; immunogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11; Conservative
                                           1 RIQRGPGRAFVT 12
                                                                     7 RIORGPGRAFYT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RIQRGPGRAFVT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7 RIORGPGRAFYT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 recognise different T
standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chong P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-550482/46.
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Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                          06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                     09-JUN-1993;
                                                                                                                                                                                                    17-OCT-2003
26-NOV-1999
                                                                                                                                                                                                                                                                                                                                                  US5951986-A.
                                                                                                                                                                                                                                                                                                                                                                            14-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Klein MH,
                                                                                                                                                                         AAY39690;
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                                                                                                                  RESULT 216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                AAY39690
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                                                                                                                                                          요 .
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/note= "This residue is bonded to the thiol sulphur of Yey(13) via a -CO-CH2- linkage, formed by introducing a bromoacetyl group onto the N-terminal and allowing the Br to condense with the Cys side chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New annular antigen scaffold cores are provided for antigens or epitopes such as HIV V3 loop peptides (e.g. the present sequence; but see also CB282818; AR890219 - AR890223), GnRH peptides, malazia antigenic peptides or bacterial capsular polysaccharides. The scaffolds comprise a ring of 3-10 Lys residues cyclised via a thioether linkage. The epitopes or antigens are bonded to each of the Lys side-chain amino groups. The Cterminus of the scaffold may be linked to a moderly such as beta-alamine or a peptide providing a T cell epitope, a lipopeptide which may provide an adjuvant effect, or another moiety providing a carrier function. The scaffolds constitute effective synthetic vaccines. The present sequence represents one of four identical thioether-cyclised HIV V3 loop peptides which are attached to each of the four Lys residues in the the annular scaffold core described in ARR90224
                                                                                                                                               /note= "this is an epsilon-Lys residue, the alpha-amino and carboxy terminals of which are incorporated into an annular antigen scaffold core of formula KKKKCX as described in AAR90224"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New poly:lysine annular core for carrying epitope(s) - esp HIV V3 loop peptide, gonadotropin releasing hormone, malarial or bacterial peptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nuclear protein antigen; systemic rheumatic disorder; human immunodeficiency virus; HIV-1; systemic lupus erythematosus; mixed connective tissue disease; scleroderma; glycoprotein 120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         epitope; autoantibody; immunoinfective cluster virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 54; DB 2;
Pred. No. 0.27;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV-1 gp120 V3 loop neutralising domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tolman RL;
                                                                                                           /note= "see above"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR62165 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 5; Page 49; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                 94GB-00020263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 70.1%;
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cunningham B, Hannah J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1995-141219/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful in vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 AA;
                                                                                                                             Modified-site
                                                                                                                                                                                                                                                                                                                                            07-0CT-1994;
                                                                                                                                                                                                                                                                                                                                                                                       15-OCT-1993;
                                                                                                                                                                                                                                                           GB2282813-A.
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25-MAR-2003
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AAR62165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-tyrosine was substd. for a val at position 11 and substrines at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-3, the Gln residue at position 3 in peptide 18 has been replaced by a Thr residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cyclic HIV PND peptide attached to annular antigen scaffold.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        72.1%; Score 55.5; DB 2; Length 14; 86.7%; Pred. No. 0.15; 1ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shirai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 annular antigen scaffold core; AASC; HIV V3 loop, lysine; principal neutralising determinant; PND; cyclic; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pendleton CD, Nara P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 33; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR90229 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                     94WO-US005142.
                                                                                                                                                                                                                                                                                                                                                                              93US-00060988
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               (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                    (USSH ) US SEC DEPT HEALTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Berzofsky JA, Ahlers JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                          HIV-1 IIIB peptide 18-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-006707/01.
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hes 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
Modified-site
           25-MAR-2003
03-AUG-1995
                                                                                                                                                                                                                                                                                                                                   13-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                            14-MAY-1993;
                                                                                                                                                                                                                                                WO9426785-A1
                                                                                                                                                                                                                                                                                           24-NOV-1994.
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                                                                                                                                                                                                       Synthetic.
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Query Match

Best Loc Matches

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Synthetic

AAR90229;

RESULT 218

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Sequence 10 AA;
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Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-0CT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-MAY-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GB2271995-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                              Scott D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW03409;
                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 221
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                                                                                                                                                                                                                                                  localised the main neutralising domains. The target of more than 80% of neutralising antibodies in HIV-1 infected sera from AIDS patients has now been found to overly the consensus binding sequence and domain A epitopes of the U1 snRNP 70% protein. In AIDS, antibody titres are too low to immunodominant targets of autoantibodies in the systemic rheumatoid disorder of mixed connective tissue disease. The titers of such autoantibodies are useful power 7. The anti-snRNP autoantibodies will cross-react with HIV-1 epitopes and are useful for treating HIV infection. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-MAR-2003 to correct PN field.) (Updated on 27-MAR-2003 to correct PN field.)
                                                                                                                                                                          Methods for treating immunoinfective cluster virus infections - utilise antibodies or fragments characteristic of auto antibodies produced by patients with rheumatic disorders.
                                                                                                                                                                                                                                          Previous immunological analyses of the V3 loop of HIV-1 (AAR62159) have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #31.
                                                                                                                                                                                                                                                                                                                                                                                                   68.8%; Score 53; DB 2; Length 10;
100.0%; Pred. No. 0.27;
ative 0; Mismatches 0; Indels
                                                                                                                                     Ehresmann G
                                                                                                                                                                                                                      Disclosure; Page 62; 106pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW76861 standard; peptide; 10 AA.
                                                                                                                (UYSC-) UNIV SOUTHERN CALIFORNIA
          Human immunodeficiency virus 1.
                                                                        94WO-US002631.
                                                                                            93US-00029850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0040581P.
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                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100.
Les 10; Conservative
                                                                                                                                    Douvas A, Takehana Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GPGRAFVTIG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                              5 GPGRAFVTIG 14
                                                                                                                                                       WPI; 1994-302689/37.
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus sp.
Homo sapiens.
                               WO9420141-A1
                                                                        10-MAR-1994;
                                                                                            11-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-FEB-1997;
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                                                   15-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW76861;
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This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents and to socines, chemotherapeutic agents and immune response modifiers. Such cortines, chemotherapeutic agents and immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, so prevents or inhibite 'bystander' apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                  New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ě
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linked to Neisseria meningitidis OMPC via a specified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                conjugate; PND; HIV; principal neutralizing determinant; OMPC; outer membrane protein complex; anionic spacer; vaccine; human immunodeficiency virus; water-soluble.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          68.8%; Score 53; DB 2; Length 10; 100.0%; Pred. No. 0.27; o. Indels ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2. .13
/note= "the peptide is cyclised"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV principal neutralizing determinant cPND495.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        anionic spacer group"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                             Claim 10; Page 119; 154pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW03409 standard; peptide; 13 AA.
(AMNA-) AMERICAN NAT RED CROSS
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Matches 10; Conservative
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                                                                          Zambidis E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RIORGPGRAF
                                                                                                                                                WPI; 1998-506315/43
                                                                                                                                                                                                                                                                                                                                       virus infection.
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Lombardo VK;

Leanza WL,

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New conjugates of outer membrane protein and HIV epitope - for generating HIV-neutralising response, have components joined by anionic spacer to ensure solubility of prod.
                                                                                                                                                   A new conjugate immunogen comprises (a) the OMPC of Neisseria meningitidis b as a protein carrier, (b) a principal neutralizing determinant (PND) of HIV as a peptidyl epitope against which immune responses are desired, and (c) a low mol. wt. anionic spacer linking and (b). The conjugate is water-soluble, yet can carry a high peptide epitope loading. It is useful as a vaccine against HIV. The present sequence is an example of a PND used in the conjugate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV-IIIB; peptide RP337; principal neutralising domain; antibodies; diagnosis; prophylaxis; therapy; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Principal neutralising domain of HIV variants - used for producing peptide(s) and antibodies for diagnosis, prophylaxis and/or therapy therapy of HIV infection.
                                                                                                                                                                                                                                                                               68.8%; Score 53; DB 2; Length 13; 100.0%; Pred. No. 0.34; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus peptide RP337,
                                                                                                                            Disclosure; Page 21; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                      AAR04441 standard; protein; 14
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89US-00407663.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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Best Local Similarity 100.
                          Marburg S,
(MERI ) MERCK & CO INC
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                                                                                                                                                                                                                                                                                                                                               3 ORGPGRAFVT 12
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Lynn DU, Petrobre J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (REPK ) REPLIGEN CORP
                                                 WPI; 1994-128412/16
                                                                                                                                                                                                                                                                                                                                   3 QRGPGRAFVT
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                                                                                                                                                                                                                                                        Sequence 13 AA;
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20-SEP-1990
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19-SEP-1989;
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                         Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
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Peptide RP335 comprises segments of the Principal Neutralising Domain (envelope protein) from isolate HIV-IIIB. The last Cys resisue is added

Claim 8 (44); Page 76; 108pp; English.

Grimaila R;

Farley J,

Javaherian K,

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Gaps

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for the purpose of crosslinking to carrier proteins. Cysteine residues can be added so that that residues at or near both ends form a disulfide bond, thus giving the peptide a loop-like configuration, which is utilised to enhance the immunogenic properties of the peptide. The peptide is capable of eliciting, and/or binding with, neutralising antibodies. The neutralising domain is bounded by cysteine residues which occur at positions 296 and 331.Peptides can be used as immunogens or screening reagents to generate or identify poly- or monoclonal Abs. See also AAR04427-R04506 and AAQ04273-Q04279. (Updated on 25-MAR-2003 to correct PR field.) (Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a T-cell epitope derived from the V3 sequence of the HIV-1 isolate HKB2, which may be linked to a B-cell epitope from the V3 (MN) loop from HIV-1. These chimeric peptides may then be used in the production of HIV-1 vaccines. These peptide sequences may also be used in the production of multimeric peptides in which the peptides are C-terminally modified by the addition of a Lys residue which is modified on its epsilon amino acid to carry an additional copy of the peptide modified on its epsilon amino acid to carry an additional copy of the peptide retreatment of AIDS by acting to displace the binding of HIV virus to human or animal cells or by disturbing the 3D organisation of the virus. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 16-OCT-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T-cell; epitope; HIV-1; core protein; p24E; B-cell; antigen; gp160; gag; pol; vaccine; multimeric peptide; AIDS; 3D organisation.
                                                                                                                                                                                                                               Revised record issued on 09-SEP-2004 : Correction to Feature Table Key
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                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                        68.8%; Score 53; DB 2; Length 14;
100.0%; Pred. No. 0.36;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cell epitope derived from V3 isolate HXB2.
                                                                                                                                                                                           on 25-MAR-2003 to correct PI field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
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(revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                       Sequence 14 AA;
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25-MAR-2003
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                                                                                                                                                                                                                                                                                                                              Local
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AAW25835;

RESULT 224

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Immunogen; vaccine; HIV-1; T-cell; B-cell; epitope; core protein; gp120;
                                                                                                                                                                                                                               HIV-1 strain HXB2 gp120 V3 loop epitope peptide.
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                           AAW67351 standard; peptide; 14 AA.
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                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
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94US-00257528.
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                                                                                                                                          (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chong P, Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-556461/47.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-JUN-1993;
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25-JAN-1999
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                                                                                       AAW67351;
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AAW67351

LD AAW6

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DT 17-(

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to new synthetic peptides comprising at least one amino acid sequence comprising an HIV gag protein T-cell epitope linked ex itse C. or N-terminus to an amino acid sequence comprising a B-cell epitope of the V3 loop of an HIV env protein, which can be used to generate vaccines against HIV-1. The T-cell epitope sequence is pref. P24M and P24H while the B-cell epitopes are derived from HIV strains including CTLB-56, V3MN, CTLB-29, CTLB-55, SF2, LAI, IIIB, RF, Z6, 2054, 174 and BXN8. The peptides are chimaeric and can be linked to a branched Lys backbone. This aequence represents the B-cell env protein V3 loop peptide from HIV-1 strain HXB2. (Updated on 25-MAR-2003 to correct PF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tandem synthetic HIV peptide(s) useful as immunogens - comprising gag
protein T-cell epitope linked to env protein B-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV; human immunodeficiency virus; gag; T-cell; B-cell; epitope; env; V3 loop; vaccine; determinant; chimaeric.
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100.0%; Pred. No. 0.36;
ive 0; Mismatches 0; Indels
                                                                                                                     Length 14;
                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV B-cell strain HXB2 env protein V3 loop peptide.
                                                                                                               DB 2;
0.36;
                                                                                                               68.8%; Score 53; DB 100.0%; Pred. No. 0.3 iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Col 21; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW25835 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Klein MH, Sia CDY, Chong P;
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(first entry)
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                                                                                                         ouery Match
Best Local Similarity 100.
Matches 10; Conservative
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                                                                                                                                                                                                                                  1 RIORGPGRAF 10
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                                                                                                                                                                                                                                                                                           5 RIORGPGRAF 14
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   standardise OS field)
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Matches 10; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 14 AA;
                                                         Sequence 14 AA;
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20-OCT-1997
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Synthetic human immunodeficiency virus-1 peptide(s) - containing T-cellepitope and B-cell epitope(s) are candidate vaccines against HV-1.
                                                                                                                     The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-cell epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitopes are generally designed based on the p24 core protein and the B-cell epitopes spream the V3 loop of the gpl20 protein from various HIV-1 epitopes from the V3 loop of the W3 loop epitope from the HIV-1 strains. This peptide represents the V3 loop epitope from the HIV-1 strain HXB2. (Updated on II-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1; human immunodeficiency virus; vaccine; T-cell epitope;
gag protein; B-cell epitope; gp41 protein; chimeric; infection.
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                                                                                                                                                                                                                                                                                                                                                                            68.8%; Score 53; DB 2; Length 14; 100.0%; Pred. No. 0.36;
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Query Match

RESULT 225

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                                                                                                                                                 The present invention describes a synthetic peptide comprising an amino acid sequence containing a T-cell epitope of an HIV gag protein linked at its C terminus to an amino acid sequence containing a B-cell epitope of an HIV gp41 protein and containing the amino acid sequence: XILKDWX2; where XI = E, A, G or Q, and X2 = A or T, or an amino acid sequence espable of eliciting an HIV-specific antiserum and recognizing the sequence XILKDWX2. The synthetic peptide is useful in vaccines against HIV infection and in diagnostic applications. AAW98882 to AAW98906, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency virus, useful antigens in producing vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV; vaccine; immunogenic composition; T cell epitope; B cell epitope;
                                                                                                                                                                                                                                                                                      Gaps
                                                                                                       Synthetic chimeric HIV polypeptides - comprising gag protein T-cell epitope linked to gp41 B-cell epitope.
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                                                                                                                                                                                                                                                                    68.8%; Score 53; DB 2; Length 14;
                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                            100.0%; Pred. No. 0.36; ive 0; Mismatches
                                                                                                                                  Example 1; Col 41-42; 41pp; English
                                                                                                                                                                                                                                                                                                                                                                 AAY39757 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                        infection; antibody; antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                              HIV1 chimeric peptide V3-HXB2.
        95US-00462507.
                                                                      Sia CDY;
                         93US-00073378
94US-00257528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Klein MH, Chong P, Sia CDY;
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94US-00257528.
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                                                   (CONN-) CONNAUGHT LAB LTD.
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                                                                                                                                                                                                                                                                           Local Similarity 100.
                                                                                                                                                                                                                                                                                                     1 RIQRGPGRAF 10
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                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
                                                                     Klein MH,
                                                                                      WPI; 1999-189590/16
                                                                                                                                                                                                                                                                                                                     5 RIORGPGRAF
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                                                                                                                                                                                                                                                  Sequence 14 AA;
        05-JUN-1995;
                         09-JUN-1993;
                                   09-JUN-1994;
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26-NOV-1999
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                                                                                                                                                                                                                                invention
                                                                     Chong P,
                                                                                                                                                                                                                                                                                                                                                                                   AAY39757;
                                                                                                                                                                                                                                                                    Query Match
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the immunogenic composition of the invention. The composition comprises a synthetic fusion polypeptide which includes a sequence encoding 1 or more T cell epitopes and a sequence encoding 1 or more B cell epitopes and a carrier. Both the T cell and B cell epitopes are derived from HIV proteins. The composition are useful as vaccines against HIV infection. The composition induces HIV-1-specific polyclonal antibodies that are opsomising and antiviral. The peptide components may be selected to induce a response against different viral isolates and in subjects who recognise different T cell epitopes. (Updated on 17-OCT-2003 to
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                                                                     This sequence represents a fragment of a HIV1 protein, and can be used in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV; LDL; low density lipoprotein; human; immune response; infection; immunodeficiency; neoplastic tissue; myalgic encephalomyelitis; MB; viral infection fatigue syndrome; tuberculosis; hepatitis; AIDS; ARC; acquired immunodeficiency syndrome; AIDS related complex; gpl20; LDL-R; HIV-infected CD4 cell; immunosuppressive peptide.
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Pred. No. 0.36;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV putative gp120 LDL-R binding region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       68.8%; Sc. 100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY22593 standard; peptide; 18 AA.
Example 1; Col 22; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus 1.
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RIORGPGRAF 10
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                                                                                                                                                                                                                                                                                                                                                                                                                         standardise OS field)
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Best Local Similarity
Local 10; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 14 AA;
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Principle neutralising domain; PND; HIV-1; gp120; envelope; vaccine;
                                                                                                                                                                             AAR26879 standard; peptide; 10 AA.
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(first entry)
                                                                                                                                                                                                                                                                                        (first entry)
                                                    9 IORGPGRAFVKL 20
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                           2 IQRGPGRAFVTI 13
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                                                                                                                                                                                                                                                                  (revised)
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                                                                                                                                                                                                                                                                                                                                    HIV epitope #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W09214489-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-FEB-1992;
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20-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                  prophylaxis
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                        AAR26879;
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                                                                                                                                      RESULT 230
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                                                                                                                                                             AAR26879
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or a mimic molecule to the cysteine rich domain of LDL receptor. The human recombinant LDL receptor forms pharmaceutical compositions for: the treatment of acquired immunodeficiency syndrome (AIDS) or ARC (AIDS related complex); reducing syncytium formation in HIV-infected CD4 cells; treating blood or body fluid or organs to neutralise/remove immunosuppressive peptides and/or viruses; or treating hepatitis A, B or C. The pharmaceutical compositions also treat a viral infection in a human or animal host. The human recombinant LDL receptor is also useful above. The human recombinant LDL receptor is a highly specific inhibitor of HIV-1 replication in vitro. (Updated on 17-OCT-2003 to standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequences given in AAR26879-93 are portions of the HIV-1 principle neutralising domain (PND) protein which were used as neutralising epitopes. HIV-1 PND is an approx. 40 amino acid region of the external envelope protein gpl20 which forms a looped structure in native gpl20. The epitopes were inserted into the BC loop of a hybrid poliovirus. This construct could be used as a vaccine. The vaccine may be used for prophylaxis or treatment of human patients infected with HIV-1. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hybrid polio virus useful as vaccine against HIV-1 infections - contains epitope of heterologous protein inserted into the BC loop of polio virus.
                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                      Length 18;
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                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                    DB 2;
                                                                                                                                                                                                                                                                                                                                  68.8%; Score 53; DB 100.0%; Pred. No. 0.4 iive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92WO-US001303.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
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Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                           1 RIQRGPGRAF 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 RIQRGPGRAF 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (REPK ) REPLIGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1992-315940/38.
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                                                                                                                                                                                                                                                                                             Sequence 18 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR26890;
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                                                                                                                                                                                                                                               field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 229
                                                                                                                                                                                                                                                                                                                                                                               Matches
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The sequences given in AAR26879-93 are portions of the HIV-1 principle neutralising domain (PND) protein which were used as neutralising epitopes. HIV-1 PND is an approx. 40 amino acid region of the external envelope protein gpl20 which forms a looped structure in native gpl20. The epitopes were inserted into the BC loop of a hybrid poliovirus. This prophylaxis or treatment of human patients infected with HIV-1. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                 Hybrid polio virus useful as vaccine against {\tt HIV-1} infections - contains epitope of heterologous protein inserted into the BC loop of polio virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Principle neutralising domain; PND; HIV-1; gp120; envelope; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       67.5%; Score 52; DB 2;
100.0%; Pred. No. 0.38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 18; 35pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR26892 standard; peptide; 10 AA.
92WO-US001303.
                                                  91US-00655669.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 . EXEXEZSX
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The peptide ellcits cytotoxic T lymphocyte (CTL) response at concns. of 10(-12) to 10(-6) M. It corresp. to residues 318-327 of HIV-1 strain IIIb gp. 160 envelope glycoprotein. It can be used for the treatment and/or prophylaxis of HIV infection. (Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent clashes with ongoing US granted patent numbers. For further information please visit the Derwent web site at www.derwent.com/dwpi/updates/MIKIS us.html.) (Updated on 25-MAR-2003 to correct PF field.) (Updated on 27-MUG-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mucosal binding compositions for generating mucosal immune response - comprises mucosal binding peptide, pref. derived from cholera toxin, and an antigen, e.g. derived from E. coli, HIV, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A novel mucosal binding compsn. (MBC) comprises a mucosal binding polypeptide linked to at least 1 antigen from a viral pathogen, which causes a sexually transmitted disease (STD), e.g. the present HIV gpl20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mucosal binding composition; mucosal binding polypeptide; antigen; viral pathogen; sexually transmitted disease; administration; vaginal; rectal; oral; immune response; secretory immunity; mucous; HIV-1; glycoprotein 120; gp120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
       Kozlowski S;
                                                                            Peptide(8) for stimulation of cytotoxic T cells specific for HIV-1 which correspond to residues 318-327 of HIV-1 gp 160 envelope
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                       Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV gp 120 antigen, component of mucosal binding compsn.
           Pendleton CD,
                                                                                                                                                                                                                                                                                                                                                                                                                   67.5%; Score 52; DB 2;
100.0%; Pred. No. 0.38;
iive 0; Mismatches
       Shirai M,
                                                                                                                                                    Disclosure; Page 8; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR95920 standard; peptide; 10 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
       Berzofsky JA, Taskeshita T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95WO-GB002708
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RGPGRAFVTI 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 RGPGRAFVTI 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (LEBE/) LEBENS M R.
(HOLM/) HOLMGREN J R.
                                           WPI; 1993-093577/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1996-268614/27.
                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10 AA;
                                                                                                                     glyco:protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-OCT-2003
14-JAN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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       à
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                   Hybrid polio virus useful as vaccine against HIV-1 infections - contains epitope of heterologous protein inserted into the BC loop of polio virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 neutralising domain (PND) protein which were used as neutralising exploses HIV-I PND is an approx. 40 amino acid region of the external envelope protein gpl20 which forms a looped structure in native gpl20. The epitopes were inserted into the BC loop of a hybrid poliovirus. This construct could be used as a vaccine. The vaccine may be used for prophylaxis or treatment of human patients infected with HIV-1. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  squence of synthetic peptide which represents immunogenic region of the loop of HIV isolate IIIb.
                                                                                                                                                                                                                                                                                                                                                                                                                             The sequences given in AAR26879-93 are portions of the HIV-1 principle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytotoxic T lymphocyte; immunogenic peptide; V3 loop; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       67.5%; Score 52; DB 2;
100.0%; Pred. No. 0.38;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (USSH ) US DEPT HEALTH & HUMAN SERVICE.
                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 23; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR33452 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1
                                                                                                                                                 92WO-US001303
                                                                                                                                                                                      91US-00655669
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                                                                                                                                                                                                                                                            Putney SD;
                                                                                                                                                                                                                       (REPK ) REPLIGEN CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 10 AA;
                                                                                                                                                                                    14-FEB-1991;
                                                                      WO9214489-A1
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prophylaxis
                                                                                                           03-SEP-1992
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17-DEC-2001
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                                                                                                                                                                                                                                                            Murray MG,
                                     Synthetic.
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Best Loca Matches

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RESULT 232

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antigen. The MBC, which is administered vaginally, rectally or orally, generates a mucosal immune response against the viral STD by allowing for the prodn. of high levels of secretory immunity, which forms the 1st line of defence against the majority of STD. (Updated on 16-OCT-2003 to standardise OS field)
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Sequence 10 AA

88888888

Length 10; 0; Indels 67.5%; Score 52; DB 2; 100.0%; Pred. No. 0.38; ative 0; Mismatches Query Match Best Local Similarity 100. 1 IQRGPGRAFV 10 2 IQRGPGRAFV 11 , g

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Gaps ö

AAW76839 standard; peptide; 10 AA. AAW76839;

25-JAN-1999 (first entry)

Pusion immunoglobulin heavy chain HIV gp120 B cell epitope #9.~

B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.

Homo sapiens. Мив вр.

WO9836087-A1.

20-AUG-1998

98WO-US002766. 13-FEB-1998; 97US-0040581P. 13-FEB-1997; (AMNA-) AMERICAN NAT RED CROSS

Zambidis E; Scott D,

WPI; 1998-506315/43.

New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency virus infection.

Claim 10; Page 119; 154pp; English.

This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, so prevents or inhibits' bypatander apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral T cell immunity 

Sequence 10 AA

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                                                                                                                                                                                                                                                                                                                                                                                             Inducing a cytotoxic T lymphocyte response - by maintaining a level of antigen in the lymphatic system of a mammal so as to provide a sustained CTL response, used to treat, e.g. AIDS.
                       Gaps
                                                                                                                                                                                  Cytotoxic T-lymphocyte response; CTL; antigen; lymphatic system; immunisation; tumour; infectious disease; immunotherapy; cancer; malignant melanoma; viral disease; hepatitis; AIDS.
                       ö
    Length 10;
                       0; Indels
    DB 2;
67.5%; brod. No. ...
100.0%; Pred. No. ...
0; Mismatches
                                                                                                                                                                 T cell epitope/MHC ligand SEQ ID NO:102
                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Page 27; 199pp, English.
                                                                                                          AAY10172 standard; peptide; 10 AA
                                                                                                                                                                                                                                                                                                                                         (CTLI-) CTL IMMUNOTHERAPIES CORP.
                                                                                                                                                                                                                                  Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                             97CA-02209815.
97US-00988320.
                                                                                                                                                                                                                                                                                           98WO-US014289.
                                                                                                                                                12-MAY-1999 (first entry)
     Query Match 67.5
Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                           Kuendig TM, Simard JJL;
                                          2 IQRGPGRAFV 11
                                                            2
                                                            1 IORGPGRAFV
                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-120514/10.
                                                                                                                                                                                                                                                                                            10-JUL-1998;
                                                                                                                                                                                                                                                                                                             10-JUL-1997;
10-DEC-1997;
                                                                                                                                                                                                                                                     WO9902183-A2
                                                                                                                                                                                                                                                                        21-JAN-1999
                                                                                                                                                                                                                           Synthetic.
                                                                                                                              AAY10172;
                                                                                          RESULT 235
                                                                                                  AAY101
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ö Gaps ö 67.5%; Score 52; DB 2; Length 10; 100.0%; Pred. No. 0.38; tive 0; Mismatches 0; Indels Query Match 67.5 Best Local Similarity 100. Matches 10; Conservative 4 RGPGRAFVTI 13 Sequence 10 AA; ઠ

given in the present invention

immunological cytotoxic T lymphocyte (CTL) response in a mammal. The method comprises: (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal and (b) maintaining the level of the antigen in the mammal's lymphatic system to maintaining the level of the antigen in the mammal's lymphatic system to maintain the immunologic CTL response. The method can be used for the delivery of e.g. a differentiation antigen, a tumour-specific multilineage antigen, an entigen, a nework of the treatment of disease such as cancer, e.g. malignam melanoma or infectious disease, e.g. viral disease such as hepatitis or AIDS. Sustained antigen delivery to the lymphatic system provides for potent CTL stimulation that takes place in the milieu of the lymphoid organ, and it sustains stimulation that is necessary to keep CTL active, cytotoxic and recirculating through the body. AAV10071 to AAV10639 represent examples of peptide antigens

The present invention describes a method of inducing and/or sustaining an

Cytotoxic T-lymphocyte response; CTL; antigen; lymphatic system; immunisation; tumour; infectious disease; immunotherapy; cancer; malignant melanoma; viral disease; hepatitis; AIDS.

Human immunodeficiency virus 1.

Synthetic

WO9902183-A2

10-JUL-1998; 10-JUL-1997; 10-DEC-1997;

21-JAN-1999

T cell epitope/MHC ligand SEQ ID NO:94.

12-MAY-1999 (first entry)

AAY10164;

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The present invention describes a method of inducing and/or sustaining an immunological cytotoxic I lymphocyte (CTL) response in a mammal. The method comprises (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal's lymphatic system to maintain the mumunological CTL response. The method can be used for the delivery of e.g. a differentiation antigen, a tumour-specific multilineage antigen, an embryonic antigen, an oncogene antigen, a mutated tumour-suppressor gene antigen, or a viral antigen. They can be used for the treatment of antigen, or a viral antigen. They can be used for the treatment of disease such as cancer, e.g. malignant melanoma or infectious disease, e.g. viral disease such as happatitis or AIDS. Sustained antigen delivery place in the militat of the lymphatic system provides for potent CTL stimulation that takes place in the militat of the lymphatic and organ, and it sustains stimulation that the militation that the militation of the lymphatic and the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     that is necessary to keep CTL active, cytotoxic and recirculating through the body. AAY10071 to AAY10639 represent examples of peptide antigens given in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Inducing a cytotoxic T lymphocyte response - by maintaining a level of antigen in the lymphatic system of a mammal so as to provide a sustained CTL response, used to treat, e.g. AIDS.
                                                                                                                                                                                                                                                                                                                                                                     Cytotoxic T-lymphocyte response; CTL; antigen; lymphatic system; immunisation; tumour; infectious disease; immunotherapy; cancer; mallgnant melanoma; viral disease; hepatitis; AIDS.
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100.0%; Pred. No. 0.38;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                             HLA Class I motif peptide SEQ ID NO:477.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 46; 199pp; English
                                                                                                                                          AAY10547 standard; peptide; 10 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus 1.
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97US-00988320
                                                                                                                                                                                                                                                           (first entry)
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     1 RGPGRAFVTI 10
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es 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10 AA;
                                                                                                                                                                                                                                                           12-MAY-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                   AAY10547;
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Matches
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Inducing a cytotoxic T lymphocyte response - by maintaining a level of antigen in the lymphatic system of a mammal so as to provide a sustained CTL response, used to treat, e.g. AIDS.

(CTLI-) CTL IMMUNOTHERAPIES CORP

Simard JJL;

Kuendig TM,

WPI; 1999-120514/10.

97CA-02209815. 97US-00988320. 98WO-US014289.

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Gaps
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                                                                                                                                                                                                                                             Score 52; DB 2; Length 10;
Pred. No. 0.38;
                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                100.08; Preu. ...
                                                                                                                                              Disclosure; Page 26; 199pp; English.
                                                                                                                                                                                                                                                                                           AAY03691 standard; peptide; 10 AA.
                                                                                                                                                                                                                          given in the present invention
                                                                                                                                                                                                                                             67.5%;
                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
Matches 10, Conservative
                                                                                                                                                                                                                                                              13
                                                                                                                                                                                                                                                                       1 RGPGRAFVTI 10
                                                                                                                                                                                                                                                                                                               (revised)
                                                                                                                                                                                                                                   Sequence 10 AA;
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07-JUN-1999
                                                                                                                                                                                                                                                                                                      AAY03691;
                                                                                                                                                                                                                                                                                    RESULT 238
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The present invention describes a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal. The method comprises: (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal is lymphatic system to maintain the immunologic CTL response. The method can be used for the delivery of e.g. a differentiation antigen, a tumour-specific multilineage antigen, an embryonic antigen, an oncogene antigen, a mutated tumour-suppressor gene artigen, or a viral antigen. They can be used for the treatment of disease such as cancer, e.g. maligiant melanoma or infectious disease, e.g. viral disease such as hepatitis or AIDS. Sustained antigen delivery con the lymphatic system provides for potent CTL stimulation that takes place in the milieu of the lymphoid organ, and it sustains stimulation the takes the body AAVIOOT to AAVIOGS represent examples of peptide antigens
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Conservative

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4 RGPGRAFVTI

1 RGPGRAFVTI 10

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AAY10164 standard; peptide; 10 AA.

RESULT 237 AAY10164 ID AAY1 XX

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CORP. Treell immune response against at least one target antigen. The kits of the invention composition, a source of one or more CD8+ T-cell (cytotoxic T lymphocytes-(TL)) epitopes of the target antigen, plus a carrier and (ii) as boosting composition, a source of one or more CD8+ T-cell (cytotoxic T lymphocytes-(TL)) epitopes of the target antigen, plus a carrier and (ii) as boosting composition a source of CTL by the source being a non-replicating or replication-impaired recombinant poxylius source being a non-replicating. If the source of CTL epitopes in (i) is a viral vector, then the vector in (ii) is from a different virus. The kits are used to generate an immune response (prophylactic or therapeutic) against pathogens or tumours, specifically against malaria cylial or parasites such as P. falciparum, or HIV, and also many other bacterial, viral or parasitic pathogens. The kits are also used for protective response against melanoma and cancer of breast or colon, and generally merever a strong CD8+ response is protective. The boosting composition may be used alone to boost a naturally primed response against malaria. The specified PVV provide an excellent booster effect, better than that protectin against sporving in complete rather than partial protection against sporving. Resulting in complete rather than partial protection against sporving. All-VSIV (H) epitope string. (Updated on 17-OCT-2003 to standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CD8+ T-cell; immune response; antigen; priming composition; CTL; epitope;
CD8+ T-cell; immune response; antigen; priming composition; CTL; epitope; cytotoxic T lymphocyte; boosting; poxvirus vector; PVV; pathogen; tumour; malaria; parasite; P. falciparum; viral; bacterial; parasite; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention relates to methods and reagents for generating a protective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Generating CD8-positive T cell response to target antigen using recombinant poxvirus - for treating or preventing malaria and HIV infection, also epitope strings from Plasmodium and HIV.
                                                                                                                                                                                                                                                                                                                                                                      Schneider J, Plebanski M;
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100.0%; Pred. No. 0.38;
tive 0; Mismatches 0; Indels
                                                                 melanoma, HIV; breast; colon; vaccination.
                                                                                                                                                                                                                                                                                                                                                                    AVS, Gilbert SC,
Blanchard T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY03655 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 43; Page 20; 85pp; English.
                                                                                                         Human immunodeficiency virus 1.
                                                                                                                                                                                                                                      98WO-GB001681.
                                                                                                                                                                                                                                                                                 97GB-00011957.
                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS INNOVATION LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1999 (first entry)
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Hanke T, Smith GL, Blan
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                                                                                                                                                                                                                                                                                   09-JUN-1997;
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Gaps

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The invention comprises (1) as priming composition, a source of one or more CD8+ T-cell immune response against at least one target antigen. The Kits catter invention comprises (1) as priming composition, a source of one or more CD8+ T-cell [cytcoxic T lymphocytes (CTL)] epitopes of the target antigen, plus a carrier and (11) as boosting composition a source of CTL epitope the same as used in (1), with this source being a non-replicating or replication-impaired recombinant poxyling vector (PVV) plus a carrier. If the source of CTL epitopes in (1) is a viral vector, then the vector in (11) is from a different virus. The kits are used to generate an immune response (proph)actic or therapeutic) against pathogens or tumours, specifically against malaria compassites such as P. falciparum, or HIV, and also many other bacterial, viral or parasite pathogens. The kits are also used for protective response against melanoma and cancer of breast or colon, and generally wherever a strong CD8+ response is protective. The boesting composition may be used alone to boost a naturally primed response against malaria. The specified PVV provide an excellent booster effect, better than that from wild-type poxvirus, resulting in complete rather than partial protection against spence of the invention wild-type virus. Sequences AAY03651-60 represent CTL peptide epitopes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to methods and reagents for generating a protective
cytotoxic T lymphocyte; boosting; poxvirus vector; PVV; pathogen; tumo
malaria; parasite; P. falciparum; viral; bacterial; parasitic; cancer;
melanoma; HIV; breast; colon; vaccination; P1A tumour antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                Generating CD8-positive T cell response to target antigen using recombinant poxvirus - for treating or preventing malaria and HIV infection, also epitope strings from Plasmodium and HIV.
                                                                                                                                                                                                                                                                                                                                            Hill AVS, Gilbert SC, Schneider J, Plebanski M; h GL, Blanchard T_{\it i}
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100.0%; Pred. No. 0.38;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV-1 CLUVAC peptide, SEQ ID NO. 16.
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                                                                                   Synthetic.
Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                 97GB-00011957.
                                                                                                                                                                                                                           98WO-GB001681.
                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS INNOVATION LTD.
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                Mcmichael AJ, Hill
Hanke T, Smith GL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10 AA;
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                                                                                                                                                                                                                                               This sequence represents a HIV-1 cluster peptide vaccine conjugate (CLUVAC) sequence. The invention relates to a method for inducing a protective mucosal cytoxic T imphocyte (CTL) response in a mammalian subject, which comprises contacting a mucosal tissue of the subject with a composition comprising a purified soluble antigen. The method can induce a protective mucosal CTL response in a subject. The method can used for protection against e.g. hepatitis A virus, papilloma virus, feline immunodeficiency virus, feline leukaemia virus, Listeria monocytogense, M. tuberculosis, M. leprae, or Giardia lamblia. The method ocr-2003 to standardise OS field)
        feline immunodeficiency virus, feline leukaemia virus, M. tuberculosis,
Listeria monocytogenes, M. leprae, Giardia lamblia,
                                                                                                                                                                                                   Method for inducing a protective mucosal cytotoxic T lymphocyte immune
                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
 protective mucosal CTL response; hepatitis A virus; papilloma virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV-1; env gene; cellular immunity; virus; therapy;
envelope glycoprotein; infection; immunisation; immune response.
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                                                                                                                                                                                                                                                                                                                                                                               Score 52; DB 2; Length 10;
Pred. No. 0.38;
                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                  Derby MA, Kelsall BL,
                                                                                                                                                                                                                                                                                                                                                                       67.5%; Sco...
100.0%; Pred. No. v.
                                                                                                                                              (USSH ) US DEPT HEALTH & HUMAN SERVICE.
                                                                                                                                                                                                                              Example 3; Page 85; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY59593 standard; peptide; 10 AA.
                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus 1,
                                                                                                                    97US-0058523P.
98US-0074894P.
                                                                                                   98WO-US019028
                                                                                                                                                                 Berzofsky JA, Belyakov IM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99EP-00401265,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-00087513.
                           immune response induction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                 4 RGPGRAFVTI 13
                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RGPGRAFVTI 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV-1 env peptide I-10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
                                                                                                                                                                                  WPI; 1999-243663/20.
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                            Sequence 10 AA;
                                                              WO9912563-A2
                                                                                                   11-SEP-1998;
                                                                                                                    11-SEP-1997;
                                                                                                                             17-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-2003
05-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-MAY-1999;
                                                                                18-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP972523-A2
                                                                                                                                                                                                               response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY59593;
                                                                                                                                                                                                                                                                                                                                                                                                Matches
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This sequence represents a fragment of the HIV-1 env protein. The invention relates to a therapeutic composition for inducing cellular invention relates to a therapeutic composition for inducing cellular immunity against a virus, which comprises a nucleic acid encoding an envelope glycoprotein. The nucleic acid may be introduced into a vector DNA or a liposome and mixed with an adjuvant to prepare a concert of into a vector DNA or a liposome and mixed with an adjuvant to prepare a virus. The therapeutic composition can be used to prevent or treat circuit prior art methods of immunishing patients against the HIV virus. The therapeutic composition can be used to prevent or treat circuit prior art methods of immunishing patients against viruses which cell turnover because of secondary responses induced by the V3 loop mutated epitopes. The full length envelope glycoprotein expressed on the cell surface or released from HIV infected cells can also trigger detrimental effects which are essential in AIDS pathogenesis. The composition provides antigen presenting cells (APCS) which contain the composition provides antigen presenting cells (APCS) which contain the composition provides antigen presenting cells (APCS) which contain the cell mediated cytotoxicity (ADCC), do not form syncytical, do not undergo apoptosis of CD4+ T cells. The composition therefore redirects immune responses towards the conserved epitope of the envelope glycoprotein, and undergo apoptosis to estandardise OS field)
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                                                                                                                                                                                                                                                            New composition for inducing viral immunity, useful for production of HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Therapeutic antigen; cytotoxic T lymphocyte; CTL; CTL immune response; cellular immune response induction method; vaccine; human; tumour; melanoma glycoprotein 75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     67.5%; Score 52; DB 3; Length 10;
100.0%; Pred. No. 0.38;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus-10 (HIV-10) peptide.
                                                                                                                                                                                                                                                                                                                                                                         Example 3; Page 11; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY67361 standard; peptide; 10 AA.
(AJIN ) AJINOMOTO CO INC. (UXJE-) UNIV JEFFERSON THOMAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RGPGRAFVTI 10
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                                                                                                                                                                                      WPI; 2000-099746/09.
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30-OCT-1998;
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                                                                                                                Kaneko Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY67361;
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RESULT 244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                             This sequence represents a human immunodeficiency virus (HIV-10) peptide used in the method of the invention. The invention relates to a method for inducing a cytocoxic T lymphocyte (CTL) immune response to non/weakly immunogenic proteins which are expressed on tumour cells. The method for inducing a cellular immune response to a non-immunogenic or weakly immunogenic target peptide expressed on tumour cells of a mammalian subject comprises administering antigen to induce a cellular immune response to the target peptide. The antigen comprises an immunogenic portion having a major histocompatibility complex (MHC) binding domain which binds to the MRC and an immune recognition domain which is a recognized by T-cells. The antigen is derived from the target peptide such that the MRC-binding portion binds to MHC with a greater affinity than the target peptide without material alteration of the immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    recognition portion. The methods are used for inducing a cellular immune response to a non-immunogenic or weakly immunogenic target peptide expressed on tumour cells of a mammalian subject. The antigens and
                                                                                                                                                                                               Induction of a cellular immune response to a weakly immunogenic protein, used to target and kill tumor cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, phage display; anti-inflammatory; antibody therapy;
inflammatory bowel disease; rheumatoid arthritis; septic shock;
multiple sclerosis; chronic inflammation; allograft rejection; panning;
tumour necrosis factor alpha; TNF; CDR3;
complementarity determining region; hybridoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 as well as polynucleotides encoding them,
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100.0%; Pred. No. 0.38;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               are used in vaccine compositions against tumour cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV peptide used to generate a mouse hybridoma.
                             (SLOK ) SLOAN KETTERING INST CANCER RES.
                                                                                   Nikolic-Zugic J, Dyall R, Houghton AN;
                                                                                                                                                                                                                                                                                      Example 2; Page 15; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY94398 standard; peptide; 10 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunogens of the invention,
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Matches 10; Conservative
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                                                                                                                                            4PI; 2000-126432/11.
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IID AAY9

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WPI; 2000-387610/33.

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The present sequence is an HIV peptide. A gene encoding a single-domain VH protein belonging to mouse VH group I(A) was cloned from a mouse VH protein belonging to mouse VH group I(A) was cloned from a mouse VH protein belonging to mouse VH group I(A) was cloned from a mouse managed by mind a generated against the present sequence in complex with H-2Dd. The gene was amplified by PCR. The 3' primer contained a sequence which randomised 9 amino acids in the third hypervariable loop (CDR3) of the VH represent generated the single-domain VH library reperfoire. CDR3 typically makes most antigen contacts in antibody combining sites. The PCR product was reamplified to avoid non-symmetric pairing of strands due PCR product was reamplified to avoid non-symmetric pairing of strands due to primer exhaustion. The final product was ligated into the phagemid to primer exhaustion. The final product was ligated into the phagemid clones capable of binding a specific antigen, e.g. Tumour necrosis factor alpha (TNFalpha), can be selected by library panning. Single-domain VH proteins can be used to treat or diagnose associated disorders. For example, disorders in which TNF plays a role include inflammatory bowel disease, rheumatoid arthritis, septic shock, multiple sclerosis, chronic inflammation and allograft rejection
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Small functional units of antibody heavy chain variable regions useful for diagnosis and treatment of disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gapa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis B virus nucleocapsid antigen; HBcAg; T cell epitope; cytolytic T lymphocyte; immunogenic; ICE; CTL; HIV; immunodominant core epitope; immunisation; mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     67.5%; Score 52; DB 3;
100.0%; Pred. No. 0.38;
iive 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY94588 standard; peptide; 10 AA.
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                                                                                Example 1; Page 18; 48pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus 1.
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(first entry)
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Best Local Similarity 100.
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 10 AA;
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10-JAN-2001
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nucleocapsid antigen (HBCAg) with a T cell epitope sequence inserted facilitated the insertion of a unique restriction site in HBCAg facilitated the insertion of the T cell epitope into the DNA encoding the immunodominant core epitope of the HBCAg. An example of a sultable insertion epitope is the present sequence, the mouse H2-d-restricted minimal ryclolytic Tlymphocyte epitope of HIV LAI gp 120. Alternatively other T cell epitopes may be inserted (AAV94584, AAV94584, AAV94585, AAV94586, AAV94587). The recombinant nucleic acid molecule may then be advantage of this method of immunisation strategies. The reagents that encode hybrid HBCAG generate an extremely high frequency cellular immune response against the CTL epitope. (Updated on 12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                           67.5%; Score 52; DB 3; Length 10;
100.0%; Pred. No. 0.38;
ive 0; Mismatches 0; Indels
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100.0%; Pro-
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|| RGPGRAFVTI 10
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Gaps

AAB15874 standard; peptide; 10 AA. AAB15874;

17-JAN-2001 (first entry)

Human chemokine derived peptide #26. 

Macrophage recruitment; chemokine derivative; MCP-1; osteoporosis; monocyte chemoattractant protein-1; inflammation; atherosclerosis; HIV; AIDS; stroke; psoriasis; autoimmune disease; hypertension; endotoxaemia; basophil-mediated disease; myocardial infarction; acute ischaemia; rheumatoid arthritis; contraception.

Synthetic.

WO200042071-A2.

20-JUL-2000

12-JAN-2000; 2000WO-US000821. 99US-00229071. 99US-00271192 12-JAN-1999; 17-MAR-1999; 01-DEC-1999;

(NEOR-) NEORX CORP.

Tatalick LM;

Grainger DJ,

WPI; 2000-499101/44.

conjugates New peptide 3, amide and heterocyclic compounds and saccharide conjuused for inhibiting chemokine induced activity and for treating e.g. stroke, vascular diseases, autoimmune diseases and tumor growth. Disclosure; Fig 18; 387pp; English

The present invention concerns the identification of a number of chemokines which can be used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences AABIS785-B15794, AABIS803-B15813 and AABIS831-B15848. These chemokine derivatives can be used to treat diseases such as autoimmune diseases, atherosclerosis, vostcoporosis, HIV infection and AIBS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotoxaemia, myocardial infarction, acute ischaemia and

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                                                                                                                                                                                                                                                               Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
rheumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used gene therapy for the same diseases, as well as in the production of
                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thibaudeau K;
                                                                       Score 52; DB 3; Length 10;
Pred. No. 0.38;
                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bridon DP, Ezrin AM, Milner PG, Holmes DL,
                                                                67.5%; Scor.
100.0%; Pred. No. v...
0; Mismatches
                                                                                                                                                                                                                                            Virus related peptide SEQ ID NO:1526.
                                                                                                                                                                                 AAB92350 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                          17-MAY-2000; 2000WO-US013576.
                                                                                                                                                                                                                                                                                                                                                                                               99US-0134406P.
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                                                                                                                                                                                                                         (first entry)
                                                                                         10; Conservative
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                                                                                                                         ||||||||||
RGPGRAFVTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-112059/12.
                                                                                Similarity
                                                  Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                   WO200069900-A2.
                              animal models
                                                                                                                                                                                                                         22-JUN-2001
                                                                                                                                                                                                                                                                                                        sapiens
                                                                                                                                                                                                                                                                                                                                                                                              17-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                       10-SEP-1999;
15-OCT-1999;
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Best Local Si
Matches 10
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AAB92350
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The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent opptides are not suitable as drug candidates as they require frequent modifying and attaching therapeutic peptides to allowin prevents or reduces the action of peptidases to increase length of activity (half infracellular uptake and interference with physiological processes. ARBO0829 to AAB9241 represent invention

Disclosure; Page 704; 733pp; English.

Sequence 10 AA;

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their corresponding DNA molecules. An immunogen comprises a portion of gag protein of HIV from an HIV clade, parts of p17 and p24, modified to prevent N-terminal myristoylation, and a synthetic polypeptide comprising human cytotoxic T-lymphocyte (CTL) epitopes of HIV protein. This immunogen is designed to elicit an HIV-specific immune response in humans. The immunogen is useful in the preparation of a medicament such as vaccine to prevent or treat HIV infection in a human subject. The invention also relates to method of stimulating anti-HIV immune response in a human subject which comprises administering one or more times an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amount of nucleic acid molecule sufficient to prime an immune response to the immunogen, or alse may be packaged within a delivery means, such as a modified vaccinia virus Ankara (WW) to boost the immune response to common portion of the immunogens. The present sequence is human immunodeficiency virus env protein derived restricted CTL epitope related to the invention. This restricted CTL epitope is presented by a murine MHC class I used for the mouse potency assay
                                                                  Human immunodeficiency virus; HIV; immunogen; anti-HIV; vaccine; gene therapy; fusion protein; modified vaccinia virus Ankara vector; MVA; cytotoxic T-lymphocyte; CTL; epitope.
                     Human immunodeficiency virus env protein derived restricted CTL epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel immunogen for stimulating anti-HIV immune response, has a portion of gag protein of HIV from HIV clade, parts of p17, p24 and synthetic polypeptide comprising human cytotoxic T-lymphocyte epitopes of HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to human immunodeficiency virus immunogens and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 52; DB 4; Length 10;
Pred. No..0.38;
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(ITAI-) INT AIDS VACCINE INITIATIVE.
(UYNA-) UNIV NAIROBI.
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14-OCT-2000; 2000GB-00025234.
                                                                                                                                                                                                                                                                                                                                   22-DEC-2000; 2000WO-GB004984.
                                                                                                                                                                             Human immunodeficiency virus
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18-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides an antigenic composition comprising an antigen with a 3-O-deacylated monophosphoryl lipid A or monophosphoryl lipid A adjuvant. The presence of the adjuvant causes an increased immune response. The antigen may be from a pathogenic bacterium, fungus, virus or parasite, a cancer cell, an allergen or from amyloid peptide protein. The composition can be used in the prevention and reatment of infection, cancer and diseases caused by amyloid deposition. It is particularly useful against HIV, Neisseria gonorrhoeae and respiratory syncytial virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antigenic composition having an antigen (e.g. viral protein) and an adjuvant, useful for enhancing humoral and cellular immune response in a host or as a prophylaxis against virus, bacterium, parasite, cancer cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                             Gaps
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antigen; infection; cancer; amyloid deposition.
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100.0%; Pred. No. 0.38;
tive 0; Mismatches 0; Indels
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                          DB 4;
0.38;
                     67.5%; Score 52; DB 100.0%; Pred. No. 0.3 Live 0; Mismatches
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                                                                                                                                                                                                                                                                                                        AAB49397 standard; peptide; 10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAY-2000; 2000WO-US013156.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-SEP-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV peptide SEQ ID NO: 12
                                                                          10; Conservative
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Les 10, Conserv
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ID AAB49397

XX AAB

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Gape

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0; Indels

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Human immunodeficiency virus type 1; HIV-1; adjuvant; immunomodulator; abba-2-macroglobulin; 3-0-deacylated monophosphoryl lipidA; MPL; GM-CSF; granulocyte macrophage colony stimulating factor; immune response; vaccine; R101 peptide.
                                                                                                                                                                                                                                      Use of 2-macroglobulin (2Masterisk), 3-0-deacylated monophosphoryl lipid A (MPL) and granulocyte macrophage colony stimulating factor (GM-CSF) for eliciting an immune response.
                                                                                                                                                                                                                                                                                    Example 2; Page 21; 53pp; English.
                                                       Human immunodeficiency virus 1.
                                                                                                                                                                                              Patel DD
                                                                                                                         27-AUG-2001; 2001WO-US026589
                                                                                                                                               25-AUG-2000; 2000US-0227624P
                                                                                                                                                                                           Haynes BF, Liao H,
                                                                                                                                                                                                                  WPI; 2002-269315/31
                                                                                                                                                                      (UYDU-) UNIV DUKE
                                                                            WO200215930-A1
                                                                                                    28-FEB-2002
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The invention relates to a composition comprising activated alpha-2-macroglobulin (alpha 2M asterisk ), 3-0-deacylated monophosphoryl lipid A (MPL) and granulocyte macrophage colony stimulating factor (GM-CSF). The invention also relates to an adjuvant suitable for use in multivalent HIV immunogenic compositions. The compositions is useful for eliciting an immune response. The present sequence is human immunodeficiency virus type 1 (HIV-1) R101 peptide used in the exemplification of the invention. (Updated on 29-AUG-2003 to standardise OS field) Gaps Length 10; Score 52; DB 5; Pred. No. 0.38; Sequence 10 AA;

ö 0; Indels Mismatches 67.5%; SCOL. 100.0%; Pre Query Match
Best Local Similarity 100.
Marches 10, Conservative

RGPGRAFVTI 10 4 RGPGRAFVTI 13

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ABG31255 standard; peptide; 10 AA. RESULT 250

(revised)
(first entry) 29-AUG-2003 21-OCT-2002 ABG31255;

GP120 classI restricted peptide.

lung cancer; infectious disease; HIV; human immunodeficiency virus; human papilloma; influenza; bacteria; parasite; vaccine; tumour cells; HSV; herpes; anti-HIV; cytostatic; immunomodulator; antibacterial; antiparasitic; cancer; lymphocytic leukaemia; lymphoma; glioblastoma; lung cancer; infectious disease; HIV; human immunodeficiency virus; gp120. ABG31255
XX
AC ABG3
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DT 29-A
DT 21-C
XX
XX
DE GP12
XX
XX
HUNG
XW HSV
X

Human immunodeficiency virus 1.

WO200256828-A2

25-JUL-2002

29-NOV-2001; 2001WO-US047808

29-NOV-2000; 2000US-0253858P

30-NOV-2000; 2000US-0250079P.

(UYRP ) UNIV ROCHESTER

Frelinger JG, Willis RA, Evans TG; Federoff HJ, Bowers WJ, Dewhurst S, Hocknell PK;

WPI; 2002-590693/63

Generating a herpesvirus amplicon particle for treating patients with cancer or infectious disease, comprises transfecting a cell with an amplicon vector, amplicon plasmid or nucleic acid sequence encoding an accessory protein.

Example 8; Page 21; 68pp; English.

This invention relates to a method for generating a herpesvirus amplicon particle comprising transfecting a cell with a Herpes simplex virus (HSV) carpiclor vector, an amplicon plasmid or a nucleic acid sequence that encodes an accessory protein. The method of the invention may have anticontent or activity. The method of the invention is useful in generating herpesvirus activity. The method of the invention is useful in generating herpesvirus amplicon particles for treating patients with cancer (e.g. chronic lymphocytic leukaemia, lymphoma, glioblastoma or lung cancer) or an infectious disease such as HIV or those caused by human papilloma virus, influenza virus, bacteria or parasite. The HSV amplicon particles or the vectors can also be useful as vaccines. Gene therapy vectors based on the vectors can also be useful as vaccines. Gene therapy vectors based on the capacity to package large amounts of genetic material (which makes them useful in expressing multiple genes or gene sequences), they have a high transduction efficiency, and they are maintained episomally, which makes them the ses prone to insertional mutagenesis. In addition to infecting many different types of cells, HSV vectors can also transduce non-replicating calls (such as tumour cells) can be removed from a patient, treated, and readministered to the patient in the course of a consistent, treated, and readministered to induce an immune response in the method of the invention. (Updated on 29-AUG-2003 to standardise OS final) 

Sequence 10 AA;

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Gaps ; 0 67.5%; Score 52; DB 5; Length 10; 100.0%; Pred. No. 0.38; Live 0; Mismatches 0; Indels 10; Conservative Query Match Best Local Similarity Matches

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Search completed: May 16, 2005, 14:40:35 Job time : 171 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Title: Perfect score:

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BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1432185 seqs, 334051727 residues Searched:

325800 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

ptodata/1/pubpaa/US10C\_PUBCOMB.pep ptodata/1/pubpaa/US10D\_PUBCOMB.pep /cgn2\_6/ptodata/1/pubpaa/US10\_NEW\_PUB.pep: /cgn2\_6/ptodata/1/pubpaa/US11\_NEW\_PUB.pep: /cgn2\_6/ptodata/1/pubpaa/US60\_NEW\_PUB.pep: /cgn2\_6/ptodata/1/pubpaa/US60\_PUBGOMB.pep: /prodata/1/puppaa/1806, NEW PUB.pp./prodata/1/puppaa/1806\_PUBGOMB.pep./prodata/1/pubpaa/1807 NEW PUB.pp./prodata/1/pubpaa/1807 NEW PUB.pep./prodata/1/pubpaa/1808 NEW PUB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./prodata/1/pubpaa/1808\_PUBGOMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep.pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pubgoMB.pep./pub ptodata/1/pubpaa/US07\_PUBCOMB.pep codata/1/pubpaa/US10A PUBCOMB codata/1/pubpaa/US10B\_PUBCOMB codata/1/pubpaa/US10C\_PUBCOMB rcgn2\_6/ptodata/1/pubpaa/US07\_PUBCOMB.p (cgn2\_6/ptodata/1/pubpaa/PCT\_NEW\_PUB.pe; (cgn2\_6/ptodata/1/mwhma./mar\_ ptodata/1/pubpaa/US09A\_PUBCOMB.| /ptodata/1/pubpaa/US09B\_PUBCOMB pubpaa/US09C\_PUBCOME pubpaa/US09\_NEW\_PUB Published\_Applications\_AA:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being pri and is derived by analysis of the total score distribution.

## SUMMARIES

|   | ;                        | Appl             | Appl             | App               | App1             | Appl             | Appl             | Appl             | | | | | | |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
|   |                          | 52,              | 49,              | 115,              | 115,              | 115,              | 115,              | 115,              | 115               | 115,              | 52,              | 52,              | 52,              | 52,              |
|   | Description              | Sequence         | Seguence         | Sequence          | Sequence          | Seguence          | Sequence          | Sequence          | Sequence          | Seguence          | Sequence         | Seguence         | Sequence         | Sequence         |
|   | QI                       | US-10-079-167-52 | US-10-360-836-49 | US-10-367-580-115 | US-10-367-593-115 | US-10-367-594-115 | US-10-367-654-115 | US-10-367-658-115 | US-10-367-668-115 | US-10-367-674-115 | US-10-653-624-52 | US-10-833-439-52 | US-10-833-745-52 | US-10-833-744-52 |
|   | 80                       | 14               | 14               | 15                | 15                | 15                | 15                | 15                | 15                | 16                | 16               | 16               | 9                | 16               |
|   | Query<br>Match Length DB | 6                | 6                | 0                 | σ                 | 6                 | 0                 | 6                 | σ                 | 0                 | 6                | 9                | 60               | σ                |
| æ | Query                    | 100.0            | 100.0            | 100.0             | 100.0             | 100.0             | 100.0             | 100.0             | 100.0             | 100.0             | 100.0            | 100.0            | 100.0            | 100.0            |
|   | Score                    | 39               | 39               | 39                | 39                | 39                | 39                | 39                | 39                | 39                | 39               | 39               | 39               | 39               |
|   | Result<br>No.            | п                | 8                | ٣                 | 4                 | S                 | 9                 | 7                 | 8                 | 6                 | 10               | 11               | 12               | 13               |

100.0%; Score 39; DB 14; Length 9; 100.0%; Pred. No. 1.3e+06;

Query Match Best Local Similarity

| Sequence 536, App Sequence 15, Appl Sequence 24, Appl Sequence 9, Appli Sequence 9, Appli Sequence 246, App Sequence 262, App Sequence 1, Appli Sequence 1, Appli Sequence 2, Appli Sequence 3, Appli Sequence 25, Appli Sequence 25 | Sequence 131, App<br>Sequence 1, Appli<br>Sequence 1, Appli<br>Sequence 315, App<br>Sequence 4, Appli<br>Sequence 10, Appli<br>Sequence 17, Appl<br>Sequence 18, Appl<br>Sequence 18, Appl<br>Sequence 18, Appl<br>Sequence 139, Appl<br>Sequence 17, Appl<br>Sequence 3, Appli<br>Sequence 6, Appli<br>Sequence 6, Appli<br>Sequence 16, Appli<br>Sequence 16, Appli<br>Sequence 7, Appli |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| US-10-239-313A-536<br>US-09-810-310-15<br>US-09-810-310-24<br>US-09-980-621-8<br>US-09-980-621-8<br>US-09-077-439A-3<br>US-10-133-210-246<br>US-10-133-210-246<br>US-10-133-210-26<br>US-10-133-210-6<br>US-10-139-313A-186<br>US-10-1239-313A-186<br>US-10-062-710-45<br>US-10-062-710-45<br>US-10-062-710-45<br>US-10-062-710-45<br>US-10-062-710-45<br>US-10-178-488-25<br>US-10-178-488-25                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |                                                                                                                                                                                                                                                                                                                                                                                            |
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| 112<br>122<br>123<br>123<br>123<br>123<br>123<br>123<br>123<br>123                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | 1220<br>120<br>120<br>100<br>100<br>100<br>100                                                                                                                                                                                                                                                                                                                                             |
| 0.0000000000000000000000000000000000000                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | 89.7<br>89.7<br>89.7<br>89.7<br>76.9<br>71.8<br>71.8<br>71.8<br>71.8                                                                                                                                                                                                                                                                                                                       |
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## ALIGNMENTS

; OTHER INFORMATION: CTL Epitope of HIV-1 gp120 US-10-079-167-52 PRIOR FILING DATE: 1999-12-09
PRIOR APPLICATION NUMBER: PCT/GB98/01681
PRIOR PILING DATE: 1998-06-09
PRIOR APPLICATION NUMBER: CFT/GB98/01681
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 1997-06-09
PRIOR FILING DATE: 2001-09-13
PRIOR FILING DATE: 2001-09-13
PRIOR FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 99
SOUTWARE: PRACESEQ for Windows Version 4.0 APPLICANT: Hill, Adrian V.S.
APPLICANT: Hill, Adrian V.S.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Reece, William
APPLICANT: Schneider, Josey
TITLE OF INVENTION: Vaccination Method
FILE REFERENCE: 2907.1000-001
CURRENT FILING DATE: 2002-02-19
PRICE APPLICATION NUMBER: US 99/454,204 Sequence 52, Application US/10079167 Publication No. US20030138454A1 ORGANISM: Unknown

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JAPELICANT: Rothman, James E.
APPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Hartl, F. Ulrich
APPLICANT: Howe, Mee, H.
APPLICANT: Houghton, Ala
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
ILIE REFERENCE: 11746/461012
CURRENT APPLICATION NUMBER: US,10/367,593
CURRENT APLICATION NUMBER: US 09/011,645
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR PILING DATE: 1996-08-16
PRIOR PILING DATE: 1995-08-18
PRIOR PILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8:0 for Windows
SEQ ID NO 115
LENGTH: 9

LENGTH: 9

LENGTH: 9

LENGTH: 9
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APPLICANT: Hartl, F. Ulrich
APPLICANT: Hoe, Mee H.
APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Todalizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Mayhew, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
TITLE OF INVENTION: WHERE: US/10/367,594
CURRENT FILING DATE: 2003-02-14
FRIOR FILING DATE: 2000-10-05
PRIOR PILING DATE: 1998-02-13
FRIOR APPLICATION NUMBER: US/10/363
FRIOR PILING DATE: 1996-08-16
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100.0%; Score 39; DB 15;
Best Local Similarity 100.0%; Pred. No. 1.3e+06;
Matches 8; Conservative 0; Mismatches 0;
                            Query Match
Best Local Similarity 100.0%; Pred. No. 1.3e+06;
Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: synthetic peptide US-10-367-593-115
                                                                                                                                                                                                                                                                                                                              Sequence 115, Application US/10367593
Publication No. US20040071721A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 115, Application US/10367594
Publication No. US20040071722A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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2 RAFVTIGK
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Sequence 1974

Sequence 1974

Publication No. US20030185854A1

GENERAL INFORMATION:

APPLICANT: Birkett, Ashley

TITLE OF INVENTION: DATICLES TO DEVELOP VACCINES AGAINST INFECTIOUS PATHOGENS:

TITLE OF INVENTION: AND WALIGNANCIES

FILE REFERENCE: 5986/1J876

CURRENT APPLICATION NUMBER: US/10/360,836

CURRENT PILING DATE: 2003-02-07

PRIOR APPLICATION NUMBER: 60/354,963

PRIOR APPLICATION NUMBER: 60/354,963

SEQ ID NOS: 86

SOFTWARE: FastSEQ for Windows Version 3.0
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JAPPLICANT: Rothman, James E.
APPLICANT: Hartl, F. Ulrich
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Maynbaw, Mark
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461061
CURRENT APPLICATION WUMBER: US 09/794, 832
FRIOR APPLICATION NUMBER: US 09/011, 645
FRIOR APPLICATION NUMBER: US 09/011, 645
FRIOR APPLICATION NUMBER: PCT/US96/13363
FRIOR APPLICATION NUMBER: PCT/US96/13363
FRIOR APPLICATION NUMBER: US 60/002, 490
FRIOR APPLICATION NUMBER: US 60/002, 490
FRIOR APPLICATION NUMBER: US 60/002, 479
FRIOR FILING DATE: 1995-08-18
FRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SROFTWARE: WordPerfect 8.0 for Windows
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Best Local Similarity 100.0%; Pred. No. 1.3e+06;
Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: human immunodeficiency virus (HIV-1)
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Sequence 115, Application US/10367580
Publication No. US20040071720A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
8; Conservative
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LENGTH: 9
Matches
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OTHER INFORMATION: synthetic peptide
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 8; Conservative
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                                                                                                                                                                                                        Length 9;
                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                        100.0%; Score 39; DB 15; 100.0%; Pred. No. 1.3e+06;
PRIOR APPLICATION NUMBER: US 60/002,490
PRIOR FILING DATE: 1995-08-18
PRIOR APPLICATION NUMBER: US 60/002,479
PRIOR FILING DATE: 1995-08-18
NUMBER OF SEQ ID NOS: 349
SOFTWARE: WordPerfect 8.0 for Windows
                                                                                                                                                           OTHER INFORMATION: synthetic peptide US-10-367-594-115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; · OTHER INFORMATION: synthetic peptide US-10-367-654-115
                                                                                                                                                                                                                                                                                                                                                                    Sequence 115, Application US/10367654
Publication No. US20040071723A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 115, Application US/10367658
Publication No. US20040071724A1
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                                                                                                                   TYPE: PRT ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 8; Conservative
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US-10-367-658-115
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GENERAL INFORMATION:

APPLICANT: Rothman, James E.

APPLICANT: Hartl, F. Ulrich

APPLICANT: Hoe, Mee H.

APPLICANT: Takechi, Yoshizumi

APPLICANT: Takechi, Yoshizumi

APPLICANT: Mayhew, Mark

APPLICANT: Mayhew, Mark

TILL OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies

FILE REFERENCE: 11746/461072

CURRENT APPLICATION NUMBER: US/10/367,668

CURRENT APPLICATION NUMBER: US 09/794,517

PRIOR APPLICATION NUMBER: US 09/794,517

PRIOR FILING DATE: 1996-08-16

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR APPLICATION NUMBER: US 60/002,479

PPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
APPLICANT: Takechi, Yoshizumi
TITLE OF INVENTION: Heat Shock Protein-Based Vaccines and Immunotherapies
FILE REFERENCE: 11746/461051
FURRENT PAPLICATION NUMBER: US/10/367,658
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 09/794,529
PRIOR APPLICATION NUMBER: US 09/71,645
PRIOR APPLICATION NUMBER: US 09/011,645
PRIOR FILING DATE: 1996-08-18
PRIOR FILING DATE: 1996-08-18
PRIOR FILING DATE: 1995-08-18
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RAFVTIGK
                                                                                                                                                                                                                                                                                                  ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Unknown
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US-10-367-674-115

US-10-367-674-115

Sequence 115, Application US/10367674

Publication No. US2040127684A1

GENERAL INPORMATION:
APPLICANT: Hartl. F. Ulxich
APPLICANT: Hartl. F. Ulxich
APPLICANT: Takenl. Yoshizumi
APPLICANT: Takenl. Yoshizumi
APPLICANT: Takenl. Yoshizumi
APPLICANT: Houghton, Alan
APPLICANT: Application NUMBER: US/10/367,674

CURRENT FILING DATE: 2003-02-14

CURRENT PILING DATE: 2003-02-14

PRIOR APPLICATION NUMBER: US 09/52,868

PRIOR PILING DATE: 2000-04-20

PRIOR PILING DATE: 1996-04-20

PRIOR APPLICATION NUMBER: US 09/011,645

PRIOR APPLICATION NUMBER: US 60/002,490

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

PRIOR FILING DATE: 1995-08-18

NUMBER OF SEQ ID NOS: 349

WUMBER WOR PELICATION NUMBER: US 60/002,490

PRIOR FILING DATE: 1995-08-18

NUMBER OF SEQ ID NOS: 349

WUMBER WOR PELICATION NUMBER: US 60/002,490

PRIOR FILING DATE: 1995-08-18

SEQ ID NO 115

LENGTH: 9

WUMBER OF SEQ ID NOS: 349

WUMBER OF SEQ ID NO 115
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                                                 Length 9;
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                                                                                            Indels
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                                            100.0%; Score 39; DB 15;
100.0%; Pred. No. 1.3e+06;
iive 0; Mismatches 0;
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APPLICANT: MCMichael, Andrew
APPLICANT: Hill, Adrian V.S.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Glabert, Jorg
APPLICANT: Plebanski, Magdalena
APPLICANT: Plebanski, Magdalena
APPLICANT: Blancher, Tomas
APPLICANT: Blanchard, Tom
TITLE OF INVENTION: Mchods and Reagents for Verint OF INVENTION: Which Generate A CD8 T Cell
FILE REFERENCE: 2907.1000-000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ) OTHER INFORMATION: synthetic peptide US-10-367-674-115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 52, Application US/10653624 Publication No. US20040131594A1 GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                        Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                    1 RAFVTIGK 8
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US-10-367-668-115
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TYPE: PRT
ORGANISM: Unknown
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LENGTH: 9
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y Sequence 52, Application US/10833744

y Publication No. US20040197349A1

y Publication No. US20040197349A1

y Publicant in Normation

y APPLICANT: McMichael, Andrew

y APPLICANT: Gilbert, Sarah C.

y APPLICANT: Gilbert, Sarah C.

y APPLICANT: Schneider, Jorg

y APPLICANT: Sarah C.

y APPLICANT: Match Geoffrey L.

y APPLICANT: Sarah C.

y APPLICANT: Match Geoffrey L.

y APPLICANT: Blanchard, Tom

y TITLE OF INVENTION: Which Generate A CD8 T Cell Immune Response

y TITLE OF INVENTION: Which Generate A CD8

y TITLE OF INVENTION: Which Canearte A CD8

y FILM DATE: 2007.1000-000

y CURRENT FILING DATE: 2003-10-16

y PRIOR PILING DATE: 1999-12-09

y PRIOR PILING DATE: 1999-12-09

y PRIOR PILING DATE: 1999-02-09

y PRIOR PILING DATE: 1999-02-09

y PRIOR PILING DATE: 1999-06-09

y PRIOR PILING DATE: 1990-06-09

                                                                                                                          GENERAL INFORMATION:
APPLICANT: Hill, Adrian V.S.
APPLICANT: Hill, Adrian V.S.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Schneider, Jorg
APPLICANT: Plebanski, Magdalena
APPLICANT: Blanchard, Tomas
APPLICANT: Blanchard, Tomas
APPLICANT: Blanchard, Tom
ITILE OF INVENTION: Methods and Reagents for Vaccination
ITILE OF INVENTION: Methods and Reagents for Vaccination
ITILE OF INVENTION: Which Generate A CD8 T Cell Immune Response
FILE REFERENCE: 2907.100-00
CURRENT APPLICATION NUMBER: US/10/686,943
PRIOR APPLICATION NUMBER: US/10/686,943
PRIOR PRILING DATE: 1999-10-09
PRIOR PILING DATE: 1999-06-09
PRIOR APPLICATION NUMBER: US/10/681
PRIOR FILING DATE: 1997-06-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 39; DB 16; Length 9; 100.0%; Pred. No. 1.3e+06; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: CTL Epitope of HIV-1 gp120
                          Sequence 52, Application US/10833745 Publication No. US20040191272A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
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ORGANISM: Unknown
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US-10-239-3134-536

US-10-239-3134-536

i Sequence 536, Application US/10239313A

i Publication No. US20030175285A1

i GENERAL INFORMATION:
    APPLICANT: KLINGUER - HAMOUR, Christine
    APPLICANT: CORVAIA, Nathalie
    APPLICANT: GOEK, Alain
    APPLICANT: GOEK, GOEK, Alain
    APPLICANT: GOEK, GO
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                                                                                                                                                                                                                                                   Gaps
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APPLICANT: Khleif, Samir N.
APPLICANT: Berzofsky, Jay A.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR CO-STIMULATION OF
TITLE OF INVENTION: METHODS AND COMPOSITIONS TO PEPTIDE ANTIGENS
FILE REFERENCE: 15280-415100US
CURRENT APPLICATION NUMBER: US/09/810,310
CURRENT PILING DATE: 2001-03-14
PRIOR PILING DATE: 2000-03-15
NUMBER OF SEQ ID NOS: 61
SOFTHARE: PALENTING VEY: 2.1
SOFTHARE: PALENTING VEY: 2.1
LENGTH: 15
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                                                                                                                                                                Length 9;
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                                                                                                                                                                Query Match
Best Local Similarity 100.0%; Pred. No. 1.3e+06;
Matches 8; Conservative 0; Mismatches 0;
; FEATURE:
; OTHER INFORMATION: CTL Epitope of HIV-1 gpl20
US-10-833-744-52
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TYPE: RTT
; ORGANISM: Human immunodeficiency virus
US-10-239-313A-536
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Patent No. US20020044948A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
PEATURE:
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Best Local Similarity 100.
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US-09-077-439A-3
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  US-09-989-621-8
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                                                                                                                                                                                                                                                                                                                                            Sequence 24, Application US/09810310
| Sequence 24, Application US/09810310
| Patent No. US20020044948A1
| GENERAL INFORMATION:
| APPLICANT: Khleif, Samir N. APPLICANT: Khleif, Samir N. TITLE OF INVENTION: IMMUNOCICAL RESPONSES TO PEPTIDE ANTIGENS
| TITLE OF INVENTION: IMMUNOCICAL RESPONSES TO PEPTIDE ANTIGENS
| FILE REFERENCE: 15280-415100US
| CURRENT APPLICATION NUMBER: US/09/810,310
| CURRENT APPLICATION NUMBER: 0201-03-14
| PRIOR APPLICATION NUMBER: 06/189,396
| NUMBER OF SEQ ID NOS: 61
| SOFTWARE: Patentin Ver: 2.1
| SEQ ID NO 24
                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
; OTHER INFORMATION: Description of Artificial Sequence: HIV PEPTIDE ; OTHER INFORMATION: ANTIGEN US-09-810-310-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: HIV PEPTIDE OTHER INFORMATION: ANTIGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cheong, Hong-Seok
TITLE OF INVENTION: Liposeomes Comprising Peptide Antigens
TITLE OF INVENTION: Liposeomes Comprising Peptide Antigens
TITLE OF INVENTION: Derived from X Protein of Hepatitis B virus
FILE REPRENCE: 0136/0E154
CURRENT APPLICATION NUMBER: US/09/989,621
CURRENT FILING DATE: 2001-11-20
PRIOR FILING DATE: 2001-11-20
PRIOR PLILING DATE: 2000-11-17
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 8
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                                                                                            Query Match
100.0%; Score 39; DB 9; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 8; Conservative 0; Mismatches 0; Indels
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| Patent No. US20020151683A1
| GENERAL INFORMATION:
| APPLICANT: Mogam Biotechnology Research Institute
| APPLICANT: Kim, Tae-Youn
| APPLICANT: Chang, Jin-Soo
| APPLICANT: Chang, Jin-Soo
| APPLICANT: Chang, Yu-Kyeong
| APPLICANT: Chang, Wa-Yoo
| APPLICANT: Choi, Myeong
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
                                                                                                                                                                                                                  1 RAFVTIGK 8
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ORGANISM: HIV
                                                                                                                                                                                                                                                                                                                             US-09-810-310-24
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Sequence 9, Application US/0982768B
Publication No. US20030165476A1
FUBLICATION NO. US20030165476A1
GENERAL INFORMATION:
APPLICANT: KINSEY, BERWA
APPLICANT: BHOGAL, BALBIR
TITLE OF INVENTION: MACROAGREGATED PROTEIN CONJUGATES AS ORAL GENETIC IMMUNIZATION DE, TITLE OF INVENTION: MACROAGREGATED PROTEIN CONJUGATES AS ORAL GENETIC IMMUNIZATION DE, TITLE OF INVENTION: MACROAGREGATED PROTEIN CONJUGATES AS ORAL GENETIC IMMUNIZATION DE, TITLE OF INVENTION: AGRNTS.
FILE REFERENCE: 201949431/10004014
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: 60/195,680
PRIOR FILING DATE: 2000-04-07
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin version 3.0
SEQ ID NOS: 13
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Publication No. US2030202989A1

GENERAL INFORMATION:
APPLICANT: Collier, R. John
APPLICANT: Blance, Steven R.
APPLICANT: Benson, Ericka L.
APPLICANT: Benson, Ericka L.
APPLICANT: Starnbach, Michael N.
TITLE OF INVENTION: Use of Toxin Peptides and/or Affinity
TITLE OF INVENTION: Handles for Delivering Compounds into Cells
FILE REFERENCE: 00246/187002
CURRENT APPLICATION NUMBER: US/09/077,439A
CURRENT FILING DATE: 1999-04-08
PRIOR FILING DATE: 1996-12-13
PRIOR PPLICATION NUMBER: US 60/019,275
PRIOR PLILOTATION NUMBER: US 60/019,275
PRIOR PLILOTATION NUMBER: US 60/008,518
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100.0%; Score 39; DB 9; Length 15; llarity 100.0%; Pred. No. 0.11; Conservative 0; Mismatches 0: Thata
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Best Local Similarity 100.
Matches 8; Conservative
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               Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                  1 RAFVTIGK
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; ORGANISM: HIV p18
US-09-827-688-9
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Query Match
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                        8 RAFVTIGK 15
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                                            1 RAFVTIGK
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US-10-239-313A-186
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                                                                                Gaps
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WS-10-133-210-246

Sequence 246, Application US/10133210

Publication No. US20030103964A1

GENERAL INFORMATION:
APPLICANT: Delisi, Charles

APPLICANT: Gulukota, Kamalakar

APPLICANT: Wacaro, Dennis

APPLICANT: Wacaro, Dennis

APPLICANT: Wang, Zhiping

APPLICANT: Wang, Zhiping

APPLICANT: Wang, Zhiping

APPLICANT: Wang, Chiping

APPLICA
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Publication No. US20030103964A1

GENERAL INFORMATION:
APPLICANT: Defisi, Charles
APPLICANT: Berzofsky, Jay
APPLICANT: Gilukota, Kamalakar
APPLICANT: Gilukota, Kamalakar
APPLICANT: Gilukota, Kamalakar
APPLICANT: Weng, Zhiping
APPLICANT: Lang, Chao
TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND
TITLE OF INVENTION: COMPOSITIONS THEREOF
TITLE OF INVENTION: WHORER: US/10/133,210
CURRENT APPLICATION NUMBER: US/10/133,210
CURRENT FILING DATE: 2002-04-26
NUMBER OF SEQ ID NOS: 281
SEQ ID NO 262

LENGTH: 15
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Query Match
100.0%; Score 39; DB 10; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 8; Conservative 0; Mismatches 0; Indels
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100.0%; Score 39; DB 14; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 8; Conservative 0; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 8; Conserv
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Sequence 186, Application US/10239313A
Sequence 186, Application WS/10239313A
Publication No. US20030175285A1
GENERAL INFORMATION:
APPLICANT: KINGUER - HAMOUR, Christine
APPLICANT: CORVAIA, Nathalie
APPLICANT: GENERAL Initiane
APPLICANT: GENERAL OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE FORM
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE FORM
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE FORM
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE FORM
CURRENT APPLICATION NUMBER: US/10/239,313A
CURRENT FILING DATE: 2002-09-19
FRIOR FILING DATE: 2000-03-23
FRIOR FILING DATE: 2000-03-23
NUMBER OF SEQ ID NOS: 697
SOFTWARE: PALENTING DATE: 2010-03-22
NUMBER OF SEQ ID NOS: 697
SOFTWARE: PALENTIN VET. 2.1
SOFTWARE: PALENTIN VET. 2.1
SOFTWARE: PALENTIN VET. 2.1
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OS-10-147910-6
SQUENCE 6, Application US/10147910
Publication No. US20030124718A1
GENERAL INFORMATION:
APPLICANT: Fuller, Deborah
APPLICANT: Fuller, James
APPLICANT: Fuller, James
APPLICANT: Shipley, Timothy
ITTLE OF INVENTION: Vaccine Composition
FILE REFERENCE: 03267-006
CURRENT APPLICATION NUMBER: US/10/147,910
CURRENT PILING DATE: 2002-05-20
FRIOR APPLICATION NUMBER: US 60/291,654
FRIOR APPLICATION NUMBER: US 60/291,655
FRIOR APPLICATION NUMBER: US 60/291,655
FRIOR RILING DATE: 2001-05-18
FRIOR FILING DATE: 2001-05-18
NUMBER OF SEQ ID NOS: 53
SOFTWARE: FasetSEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 15
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GENERAL INFORMATION:

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                                                                                                  Sequence 2, Application US/10787880

Sequence 2, Application US/1078781

Sequence 2, Application No. US2050025777A1

GENERAL INFORMATION:

APPLICANT: Pohlmann, Edward L.

APPLICANT: Sheeby, Michael J.

APPLICANT: Barton, Kenneth A.

TITLE OF INVENTION: PARTICLE-WEDIATED DELIVERY OF ANTIGENS
FILE REFERENCE: 03257-018

CURRENT APPLICATION NUMBER: US/10/787,880

CURRENT FILING DATE: 2004-02-27

PRIOR APPLICATION NUMBER: US/9/191,772

PRIOR APPLICATION NUMBER: US/9/191,772

PRIOR PILING DATE: 1998-11-13

NUMBER OF SEQ ID NOS: 2

SOFTWARE: FRRESEQ for Windows Version 4.0

LENGTH: 15
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i Sequence 44, Application US/10062710

i Publication No. US20030049253A1

i GENERAL INFORMATION:

APPLICANT: Li, Frank Q.

APPLICANT: Chu, Yong-Liang

APPLICANT: Qiu, Jian-Tai

TITLE OF INVENITION: Polymeric Conjugates for Delivery of

TITLE OF INVENITION: Polymeric Conjugates for Delivery of

TITLE OF INVENITION: Wia Peptide Vaccines

TITLE OF INVENITION: Wia Peptide Vaccines

TITLE OF INVENITION: Wia Peptide Vaccines

TITLE OF INVENITION: 2002-02-05

CURRENT APPLICATION NUMBER: US/10/062,710

CURRENT FILING DATE: 2001-08-08

NUMBER OF SEQ ID NOS: 232

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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; Sequence 45, Application US/10062710
; Publication No. US20030049253A1
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Best Local Similarity
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8 RAFVTIGK 15
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RAFVTIGK 14
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; ORGANISM: HIVGD120
US-10-787-880-2
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Sequence 3, Application US/09813659

Sequence 3, Application US/09813659

Patent No. US2002001298941

GENERAL INFORMATION:

APPLICANT: Ledbetter.

APPLICANT: Lineley, Peter S.

APPLICANT: Lineley, Peter S.

APPLICANT: Gilliland, Lias K.

TITLE OF INVENTION:

TITLE OF INVENTION: AND METHODS OF PRODUCING BISPECIFIC FUSION PROTEINS

TITLE OF INVENTION: PUSION PROTEINS IN A MAMMALIAN CELL

TITLE OF INVENTION: PUSION PROTEINS IN A MAMMALIAN CELL

TITLE OF INVENTION: PUSION PROTEINS IN A MAMMALIAN CELL

TITLE OF INVENTION: PUSION PROTEINS IN A MAMMALIAN CELL

TITLE OF INVENTION: PUSION PROTEINS IN A MAMMALIAN CELL

TITLE OF INVENTION: PUSION PROTEINS IN A MAMMALIAN CELL

TITLE OF INVENTION NUMBER: US/09/813,659

CURRENT APPLICATION NUMBER: 08/549,067

PRIOR FILING DATE: 1995-10-05

PRIOR PILING DATE: 1993-02-01

NUMBER OF SEQ ID NOS: 32

SOFTWARE: PATENTIN VET. 20

TWENT PRIOR PILING DATE: 1993-02-01

NUMBER OF SEQ ID NOS: 32

SEQ ID NO 3

LUMBER OF PATENTIN PRIOR

TWENT PRIOR PILING DATE: 1993-02-01

NUMBER OF PATENTIN PRIOR

PRIOR PILING DATE: 1993-02-01

NUMBER OF PATENTIN PRIOR

PRIOR PILING DATE: 1993-02-01

NUMBER OF PATENTIN PRIOR

PRIOR PILING DATE: 1993-02-01

NUMBER OF PATENTIN PRIOR

PRIOR PILING DATE: 1993-02-01

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PRIOR PILING DATE: 1993-02-01

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PRIOR PILING DATE: 1993-02-01

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PRIOR PILING DATE: 1993-02-01

NUMBER OF PATENTIN PRIOR

PRIOR PILING DATE: 1993-02-01

NUMBER OF PATENTIN PRIOR

TYPE PRIOR

PRIOR PILING DATE: 1993-02-01

NUMBER OF PATENTIN PRIOR

PRIOR PILING DATE: 1993-02-01
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Best Local Similarity 100.0%; Pred. No. 0.14;
Matches 8; Conservative 0; Mismatches 0; Indels
APPLICANT: Chu, Yong-Liang
APPLICANT: Chu, Jian-Tai
TITLE OF INVENTION: Polymeric Conjugates for Delivery of
TITLE OF INVENTION: MHC-Recognized Epitopes
TITLE OF INVENTION: Wia Peptide Vaccines
FILE REFERENCE: 3781-001-27
CURRENT APPLICATION NUMBER: US/10/062,710
CURRENT FILING DATE: 2002-02-05
PRIOR APPLICATION NUMBER: US 60/310,498
PRIOR FILING DATE: 2001-08-08
NUMBER OF SEQ ID NOS: 232
SOFTWARE: PREESEQ for Windows Version 4.0
SEQ ID NO 45
LENGTH: 18
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, OTHER INFORMATION: HIV Helper-T Cell Epitopes
US-10-062-710-45
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 8; Conservative
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US-09-813-659-3
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15 RAFVTIGK 22
1 RAFVTIGK
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                   US=10-143-51044-3

| Cards-chub.
| Sequence 3, Application US/10283610A
| Publication No. US2003019876A1
| GENERAL INFORMATION:
| APPLICANT: Ledbetter, Jeffrey A. APPLICANT: Linaley, Peter S. APPLICANT: Linaley, Peter S. APPLICANT: Linaley, Peter S. APPLICANT: Gilliand, Lisa K. TITLE OF INVENTION: EXPRESSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
| TITLE OF INVENTION: AND METHODS OF PRODUCING BISPECIFIC FUSION PROTEINS
| TITLE OF INVENTION: EXPRESSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
| TITLE OF INVENTION: AND METHODS OF PRODUCING BISPECIFIC FUSION PROPER PROPERTY APPLICATION NUMBER: US/10/283,610A
| CURRENT FILING DATE: 2002-10-29
| PRIOR PILING DATE: 1995-10-05
| PRIOR PILING DATE: 1995-10-05
| PRIOR PILING DATE: 1993-09-13
| PRIOR PILING DATE: 1993-09-13
| PRIOR PILING DATE: 1993-09-13
| PRIOR PILING DATE: 1993-02-01
| WUMBER OF SEQ ID NOS: 35
| SEQ ID NO 3
| LEASTHACE PRICETION NUMBER: OB/121,054
| PRIOR PILING DATE: 1993-02-01
| WUMBER OF SEQ ID NOS: 35
| SEQ ID NO 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 39; DB 14; Length 21; Best Local Similarity 100.0%; Pred. No. 0.15; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 39; DB 15; Length 20; 100.0%; Pred. No. 0.14;
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-hag 0; Indels
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ORGANISM: Unknown Organism
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Best Local Similarity 100.
Matches 8; Conservative
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ORGANISM: Homo sapiens
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                 US-10-283-610A-3
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GENERAL INFOGRANTION: TO IMPURIOR APPLICANT: DE LOYS, ROBERT

TITLE OF INVENTION: TO IMMUNICOGICALLY IMPORTANT EPITORES AND THEIR USE IN

TITLE OF INVENTION: TO IMMUNICOGICALLY IMPORTANT EPITORES OF

TITLE OF INVENTION: A PROCESS FOR DETERMINATION OF ANTIBODIES OF

TITLE OF INVENTION: BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT

TITLE OF INVENTION: BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT

TITLE OF INVENTION: CONTAINING THEM

FILE REFERENCE: 2752-11

CURRENT PELLING DATE: 2030-07-18

PRIOR APPLICATION NUMBER: US/09/576,824A

PRIOR APPLICATION NUMBER: 08/723,425

PRIOR APPLICATION NUMBER: 09/146,028

PRIOR PILING DATE: 1996-09-30

PRIOR PILING DATE: 1993-03-08

PRIOR PILING DATE: 1993-03-08

PRIOR PILING DATE: 1993-03-08

PRIOR FILING DATE: 1993-03-08

PRIOR PILING DATE: 1993-03-08

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; Sequence 131. Application US/10777053
; Publication No. US20040132088A1
; GENERAL INFORMATION:
; APPLICANT: Simard, John J. L.
; APPLICANT: Olu, Zhiyong
; APPLICANT: Olu, Zhiyong
; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING EPITOPES OF
TITLE OF INVENTION: EXPRESSION VECTORS AND METHODS FOR THEIR DESIGN
; TITLE OF INVENTION: TARGET-ASSOCIATED ANTIGENS AND METHODS FOR THEIR DESIGN
; FILE REFERENCE: MANNK. 02201
; CURRENT PAPPLICATION NUMBER: 10/292,413
; PRIOR FILING DATE: 2004-02-10;
; PRIOR FILING DATE: 2001-11.07
; PRIOR FILING DATE: 2001-11.07
; PRIOR FILING DATE: 2001-11.07
; NUMBER OF SEQ ID NOS: 979
; SOFTWARE: PSESSEQ for Windows Version 4.0
; SEQ ID NO 131
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100.0%; Score 39; DB 17;
Best Local Similarity 100.0%; Pred. No. 0.18;
Matches 8; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Human Immunodeficiency Virus
Sequence 160, Application US/10621675
Publication No. US20050049398A1
GENERAL INFORMATION:
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APPLICANT: SHIBA, KIYOTAKA
TITLE OF INVENTION: MULTIFUNCTIONAL BASE SEQUENCE AND ARTIFICIAL GENE CONTAINING THE 5.
FILE REFERENCE: 4439-4004
                                                                                                                                                                APPLICANT: CORVAIA, Nathalie
APPLICANT: CORVAIA, Nathalie
APPLICANT: CORVAIA, Nathalie
APPLICANT: CORVAIA, Nathalie
APPLICANT: CORVAIA, Nathalie
APPLICANT: BECK. Alain
APPLICANT: GOBTSCH, Liliane
TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS
TITLE OF INVENTION: NO F APYSIOLOGICALLY ACCEPTABLE STRONG ACID
TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID
FILE REFERENCE: 343 727 - US
CURRENT PILING DATE: 2002-09-19
PRIOR APPLICATION NUMBER: PCT 01/70772
SOFTWARE: PALENT PLING DATE: 2001-03-22
NUMBER OF SEQ ID NOS: 697
SOFTWARE: PALENT PLING DATE: 2010-03-22
NUMBER OF SEQ ID NOS: 697
SEQ ID NO 535
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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100.0%; Pred. No. 1.3e+06;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/311,111
CURRENT FILING DATE: 2002-12-13
PRIOR APPLICATION NUMBER: UP 2000-180997
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin version 3.1
SEQ ID NO 4
LENGTH: 7
                                                                                                    Sequence 535, Application US/10239313A Publication No. US20030175285A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
; OTHER INFORMATION: Designed peptide
US-10-311-111-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/10311111 Publication No. US20030121065A1 GENERAL INFORMATION:
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; Sequence 4, Application US/10398932
; Publication No. US20040171803A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 7; Conservative
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Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
US-10-239-313A-535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 RAFVTIG 12
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ORGANISM: artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RAFVTIG 7
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                                                                                                                                                                                                                            Sequence 1, Application US/10311111
Publication No. US20030121065A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: WULTIFUNCTIONAL BASE SEQUENCE AND ARTIFICIAL GENE CONTAINING THE FILE REFERENCE: 4439-4004
CURRENT APPLICATION WUMBER: US 10/10/311,111
CURRENT FILING DATE: 2000-180997
PRIOR APPLICATION NUMBER: UP 2000-180997
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin Version 3.1
SEQ ID NO 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Description of Artificial Sequence: Synthetically Designed; OTHER INFORMATION: Peptide
US-10-398-932-1
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                              Gaps
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US-10-398-932-1
i Sequence 1, Application US/10398932
i Sequence 1, Application US/10398932
i Fublication No. US20040171803A1
i GENERAL INFORMATION:
APPLICANT: SHIBA, KIYOTAKA
APPLICANT: OHNO, TSUMEN
ITILE OF INVENTION: OF EPITOPE
CURRENT PELLOATION NUMBER: US/10/398,932
CURRENT FILING DATE: 2003-04-11
CURRENT PELLOATION NUMBER: PT/JP01/08893
PRIOR FILING DATE: 2001-10-10
PRIOR FILING DATE: 2001-10-13
NUMBER OF SEQ ID NOS: 55
SOFTWARE: PALCHING OF SEQ ID NOS: 55
SEQ ID NO 1
MANDER OF SEQ ID NOS: 21
SEQ ID NO 1
MANDER OF DATE: 2001-10-13
SEQ ID NO 1
MANDER OF SEQ ID NOS: 21
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Pred. No. 1.1;
0; Mismatches 1; Indels
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  87.5%; Pred. No. 1.3e+06;
tive 1; Mismatches 0; Indels
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CTHER INFORMATION: Designed peptide

US-10-311-111-1
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87.5%;
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Best Local Similarity 87.5%;
Matches 7; Conservative
Similarity 87.5
7; Conservative
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                                                                   1 RAFVTIGK 8
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2 RAFVTLGK 9
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Best Local Similarity
Matches 7; Conserva
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12 RTFVTIGK 19
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ORGANISM: artificial
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Best Local
Matches
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APPLICANT: Bowers, William J.
APPLICANT: Bowers, William J.
APPLICANT: Bowers, William J.
APPLICANT: Bowers, William J.
APPLICANT: Fralinger, John G.
APPLICANT: Fralinger, John G.
APPLICANT: Evans, Thomas J.
APPLICANT: Debuhurst, Stephen
APPLICANT: Debuhurst, Stephen
APPLICANT: Powenblatt, Joseph D.
ITILE OF INVENTION: HELPER VIRUS-FREE HERPESVIRUS AMPLICON
ITILE OF INVENTION: PARTICLES AND USES THEREOF
FILE REFERENCE: 12610-011001
CURRENT APPLICATION NUMBER: US/09/997,848A
CURRENT FILING DATE: 2000-110
FRIOR APPLICATION NUMBER: US 60/253,858
FRIOR FILING DATE: 2000-11-29
FRIOR FILING DATE: 2000-11-29
FRIOR FILING DATE: 2000-11-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Dewhurst, Stephen
APPLICANT: Tolba, Khaled A.
APPLICANT: Tolba, Khaled A.
APPLICANT: Tolba, Khaled A.
APPLICANT: Tolba, Khaled A.
APPLICANT: Rosenblatt, Joseph D.
TITLE OF INVENTION: HELPER VIRUS-FREE HERPESVIRUS AMPLICON
TITLE OF INVENTION: PARTICLES AND USES THEREOF
FILE REPERBENCE: 12610-011001
CURRENT APPLICATION NUMBER: US 60/253,858
PRIOR APPLICATION NUMBER: US 60/253,858
PRIOR FILING DATE: 2000-11-29
PRIOR FILING DATE: 2000-11-29
PRIOR FILING DATE: 2000-11-30
NUMBER OF SEQ ID NOS: 18
SOFTWARE: PRAESEQ for Windows Version 4.0
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1.3e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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APPLICANT: Bowers, William J.
APPLICANT: Frelinger, John G.
APPLICANT: Willis, Richard A.
APPLICANT: Evans, Thomas J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
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; OTHER INFORMATION: Peptide
US:10-398-932-4
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APPLICANT: Callebaut, Christian
APPLICANT: Callebaut, Christian
APPLICANT: Callebaut, Christian
APPLICANT: Krust, Bernard
APPLICANT: Briand, Gannea
APPLICANT: Briand, Gann-Paul
APPLICANT: Guichard, Giles
APPLICANT: Guichard, Giles
TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC USES.
TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC USES.
CURRENT APPLICATION NUMBER: US/09/825,886
CURRENT FILING DATE: 2001-07-26
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                                                                          TITLE OF INVENTION: JOURNAL
TITLE OF INVENTION: OF EPITOPE
CURRENT APPLICATION UNMER: US/10/398,932
CURRENT FILING DATE: 2003-04-11
PRIOR FILING DATE: 2001-10-10
PRIOR FILING DATE: 2001-10-10
PRIOR FILING DATE: 2001-10-13
PRIOR FILING DATE: 2000-10-13
NUMBER OF SEQ ID NOS: 55
SOFTWARE: PATENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 7; Score 30; DB 16; Length 7; Best Local Similarity 100.0%; Pred. No. 1.3e+06; Matches 6; Conservative 0; Mismatches 0; Indels
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PRIOR FILING DATE: 1999-09-10
PRIOR PILING DATE: 1999-09-10
PRIOR PILING DATE: 1998-00-12
PRIOR PILING DATE: 1998-01-12
PRIOR APPLICATION NUMBER: 60/040,969
PRIOR FILING DATE: 1997-03-12
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 30
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; Sequence 30, Application US/09825886
; Publication No. US20020076693A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
FBATURE:
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Matches 6; Conserv
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GENERAL INFORMATION
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APPLICANT: Diamond, David C.
APPLICANT: Diamond, David C.
APPLICANT: Diamond, David C.
APPLICANT: Diamond, David C.
APPLICANT: Lei, Xiang-Dong
TITLE OF INVENTION: EXPRESSION VECTORS ENCODING EPITOPES OF
TITLE OF INVENTION: TARGET-ASSOCIATED ANTIGENS AND METHODS FOR THEIR DESIGN
FILE REPERENCE: MANNK: 0.22-10
CURRENT APPLICATION NUMBER: US/10/77,053
CURRENT PILING DATE: 2004-02-10
PRIOR APPLICATION NUMBER: 60/326,413
PRIOR PELLING DATE: 2001-11-07
PRIOR PELLING DATE: 2001-11-07
PRIOR PELLING DATE: 2001-11-07
SEQ ID NOS: 979
SEQ ID NOS: 979
LENGTH: 9
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APPLICANT: Fantini, Jacques
TITLE OF INVENTION: HIVGF120-INDUCED BOB/GPR15 ACTIVATION
FILE REFERENCE: 21101.00221
CURRENT APPLICATION NUMBER: US/10/494,161
CURRENT FILING DATE: 2004-04-29
PRIOR APPLICATION NUMBER: PCT/US 02/34336
PRIOR PILING DATE: 2002-10-25
PRIOR FILING DATE: 2001-10-29
NUMBER OF SEQ ID NOS: 17
SOFTWARR: FastSEQ for Windows Version 4.0
SEQ ID NO 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Description of Artificial Sequence:/note; OTHER INFORMATION: Synthetic Construct
US-10-494-161-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               71.8%; Score 28; DB 17; Length 9; 100.0%; Pred. No. 1.3e+06; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Human Immunodeficiency Virus
US-10-777-053-139
                                                                                                                           Sequence 139, Application US/10777053
Publication No. US20040132088A1
GENERAL INFORMATION:
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Sequence 17, Application US/10494161

Publication No. US20050042747A1

GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity luv.
6, Conservative
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Best Local Similarity 100.
Matches 6; Conservative
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RAFVTI 9
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Sequence 3, Application US/09858349
Patent No. US20020012909A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: SMALL FUNCTIONAL UNITS OF ANTIBODY HEAVY CHAIN VARIABLE REGIONS
FILE REPERENCE: 97534-2800
CURRENT APPLICATION NUMBER: US/09/858,349
CURRENT FILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.1
SEQ ID NO 3.1
SEQ ID NO 3.1
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US-09-810-310-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Khief, Samir N.
APPLICANT: Berzofsky, Jay A.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR CO-STIMULATION OF
TITLE OF INVENTION: IMMUNCACICAL RESPONSES TO PEPTIDE ANTIGENS
FILE REFERENCE: 15280-415100US
CURRENT APPLICATION NUMBER: US/09/810,310
CURRENT APPLICATION NUMBER: 60/189,396
PRIOR APPLICAND NUMBER: 60/189,396
PRIOR FILING DATE: 2000-03-15
                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: mouse hybridoma specific for H-2D + RGPGRAFVTI peptide US-09-858-349-3
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APPLICANT: Cho, Hearn Jay
APPLICANT: Richman, Douglas
APPLICANT: Horner, Anthony A.
ITILE OF INVENTION: Method for Increasing a Cytotoxic T
                                                                                                                                                                                                                                                                                                                                                                                                          71.8%; Score 28; DB 9;
100.0%; Pred. No. 19;
iive 0; Mismatches (
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ORGANISM: Artificial Sequence
FEATURE:
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 16
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 71.8
Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity 100.
Matches 6; Conservative
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JS-09-858-349-3
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 45
US-09-087-513-7

Sequence 7, Application US/09087513

Publication No. US20020182180A1

GENERAL INFORMATION:

APPLICANT: KOADEKO, YUTARO

TITLE OF INVENTION: METHOD OF INDUCING IMMUNITY TO VIRUSES

PILE REFERENCE: 0010-0929-0X

CURRENT APPLICATION NUMBER: US/09/087,513

CURRENT FILING DATE: 1998-05-29

NUMBER OF SEQ ID NOS: 23

SEQ ID NO 7

LENGTH: 10
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100.0%; Pred. No. 19;
cive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: HIV-1 class I-restricted gp120 peptide US-09-820-484-8
FILE REPERENCE: 06510-188US1
CURRENT APPLICATION NUMBER: US/09/820,484
CURRENT FILING DATE: 2001-03-28
FRIOR PILING DATE: 2000-03-28
FRIOR PILING DATE: 2000-05-11
PRIOR PILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 8
SEQ ID NO 8
LENGTH: 10
TYPE: PRI
ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity 100.
Matches 6; Conservative
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Search completed: May 16, 2005, 08:13:00 Job time : 132 secs

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OM protein - protein search, using sw model

May 16, 2005, 09:40:42; Search time 66 Seconds (without alignments) 46.880 Million cell updates/sec

Run on:

1 rafvtigk 8 Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

2105692 segs, 386760381 residues Searched:

768190 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 250 summaries

Database :

A\_Geneseq\_16Dec04:\*
1: geneseqp1980s:\*
2: geneseqp1990s:\*
3: geneseqp2000s:\*
4: geneseqp2001s:\*
5: geneseqp2001s:\*
6: geneseqp2001ss:\*
7: geneseqp2003bs:\*
8: geneseqp2004ss:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

|           |            | Description | Aar38167 V3 loop p | Aar72313 Anti-HIV | HIV      | Aar46523 HIV gp120 | Aar62144 HIV-1 gpl | -        | Aay03692 Amino aci | _        | ZHC<br>ZHC | Ade79993 HIV1 carr | Adk50933 Breast/bl | Adr69467 Novel hyb | Aaw76840 Fusion im | H        | Aaw34472 HIV gp120 | Aar62152 HIV-1 gpl | HIV 9    | _        | Aaw62890 Peptide s | -        | Aar33336 Sequence | ¤        | HIV-1    | HIV-1    | Aaw09264 HIV-1 Btr |
|-----------|------------|-------------|--------------------|-------------------|----------|--------------------|--------------------|----------|--------------------|----------|------------|--------------------|--------------------|--------------------|--------------------|----------|--------------------|--------------------|----------|----------|--------------------|----------|-------------------|----------|----------|----------|--------------------|
| SUMMARIES |            | ΩI          | AAR38167           | AAR72313          | AAB68603 | AAR46523           | AAR62144           | AAY10157 | AAY03692           | AAU96033 | ABG79839   | ADE79993           | ADK50933           | ADR69467           | AAW76840           | AAW19056 | AAW34472           | AAR62152           | AAW54932 | AAW22327 | AAW62890           | AAM99433 | AAR33336          | AAR48604 | AAR66416 | AAR66417 | AAW09264           |
|           |            | 80          | 7                  | 7                 | 4        | N                  | ~                  | 7        | ~                  | ហ        | Ŋ          | 7                  | æ                  | œ                  | N                  | ~        | ~                  | N                  | N        | N        | N                  | 4        | ~                 | ~        | 7        | ~        | ~                  |
|           |            | Length DB   | 80                 | σ.                | œ        | 6                  | σ                  | 6        | 60                 | 6        | D          | σ                  | σ                  | თ                  | 10                 | 11       | 11                 | 12                 | 12       | 13       | 13                 | 13       | 14                | 14       | 14       | 14       | 14                 |
| ,         | *<br>Query | Match       | 100.0              | 100.0             | 100.0    | 100.0              | 100.0              | 100.0    | 100.0              | 100.0    | 100.0      | 100.0              | 100.0              | 100.0              | 100.0              | 100.0    | 100.0              | 100.0              | 100.0    | 100.0    | 100.0              | 100.0    | 100.0             | 100.0    | 100.0    | 100.0    | 100.0              |
|           |            | Score       | 39                 |                   |          | 39                 | 39                 | 39       | 39                 | 39       | 39         |                    | 39                 |                    |                    | 39       | 39                 | 39                 | 39       | 39       | 39                 | 39       | 39                | 39       | 39       | 39       | 39                 |
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| HIV gp120 Antigenic Antigenic Arceptor Fusion im HIV-1 iso HIV-1 iso HIV-1 iso HIV-1 iso HIV-1 iso Beptide P Peptide P Peptide P HIV-1 str Sequence Sequence Sequence Sequence Heterocon HiV-1 gp1 HIV-1 gp1 HIV-1 pp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | le 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |
| HIV gpl20 Antigenic Antigenic Antigenic Anceptor Fusion im Fusion im HIV-1 iso Human imm Peptide Peptide Peptide Peptide Gpl20 V3 HIV-1 str                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | Parision of the state of the st                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
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| Aaw54930<br>Aaw34475<br>Aaw34475<br>Aaw349180<br>Aaw76901<br>Aaw76476<br>Aau060522<br>Aar42153<br>Aar42153<br>Aar42153<br>Aar42153<br>Aar64016<br>Aar64016<br>Aar64016<br>Aar64016<br>Aar26665<br>Aar26670<br>Aar26670<br>Aar38190<br>Aar38190<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100<br>Aar38100                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | 10000000000000000000000000000000000000                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
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AAR72313 standard; peptide; 8 AA

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HIV gpl20 V3. loop-derived peptides (AAR38170-87) are successful in generating CTL responses, esp. peptide R15K (AAR38187); the T-helper cell-inducing peptide includes the sequence C19A (AAR38164); HIV infection-inhibiting peptides are given in AAR38188-206, and are esp. peptides R15K, N24G, E13V, R8K, T13O and H13N (AAR38165-69). The peptides may also be derived from an influenza virue protein or a sendai virus protein (AAR41014-15). It was observed that peptide R8K (amino acids 322-323), with sequences derived from the V3 loop of HIV-1 IIIB, inhibits HIV-1 mifection of primary human T cells by 66% at 1 microg/ml (ca. 1.25 microw). (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG -2003 to correct D8 field.)
Abp60029 HIV antig
Abr39122 HIV-1 gp1
Abp72314 HIV-1 p18
Ada50228 Human imm
Ade79992 HYU carr
Ade79994 HIV1 carr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide composition for treating and preventing viral infections - comprise CTL-inducing epitope and HIV infection-inhibiting sequence or T helper cell-inducing sequence.
                                                                                                                                                                                                                                                                                                                     gp120; HIV-1; cytotoxic T-lymphocyte; CTL; T-helper; AIDS; infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 39; DB 2; Length 8; 100.0%; Pred. No. 1.8e+06; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Arlinghaus RB, Platsoucas CD, Nehete PN;
                                                                                                           ALIGNMENTS
 ABP60029
ABR39122
ABP72314
ADA50228
ADE79992
ADE79994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 19; Page 95; 130pp; English.
                                                                                                                                                                                AAR38167 standard; peptide; 8 AA
                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              91US-00800932
92US-00945865
                                                                                                                                                                                                                                                                                                                                                                                                                                    92WO-US010378
                                                                                                                                                                                                                                     (revised)
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    22222
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                                                                                                                                                                                                                                                                                           V3 loop peptide R8K.
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Best Local Similarity
  71.8
71.8
71.8
71.8
71.8
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                                                                                                                                                                                                                                                                                                                                                                                                                                    02-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-DEC-1991;
16-SEP-1992;
                                                                                                                                                                                                                                    27-AUG-2003
25-MAR-2003
12-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                             WO9310816-A1
                                                                                                                                                                                                                                                                                                                                                                                                         10-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sastry JK,
  AAR38167;
                                                                                                                                                    RESULT 1
AAR38167
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Benjouad A, Yahi N, Penouillet E, Mabrouk K; Van Rietschoten J, Rochat H;

93GB-00018901.

MCKELVEY I E.

(MCKE/)

Sabatier JM, Gluckman J,

ARME-) ARMEL SA

15-JUN-1994;

94WO-GB001992

13-SEP-1994; 13-SEP-1993;

23-MAR-1995

WO9507929-A1

Synthetic.

Multiple branch peptide construction; MBPC; HIV-1; human immunodeficiency virus type 1; virucide.

Anti-HIV MBPC peptide moiety.

(revised)
(first entry)

25-MAR-2003 20-OCT-1995

AAR72313;

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                                                                                                                                                                                                                                                                                                                                                                       Multiple branch peptide constructions (MBPCs) are formed from the V3 loop of HIV-1 gpl20. Each MBPC includes multiple peptide moieties incorporating the GPGR consensus sequence, each attached to the amino group of a lysine residue, forming a dendritic structure. A peptide moiety (AAR72313) not including the consensus was unable to inhibit HIV-1 syncytia formation. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                        Multiple branch peptide constructions formed from the V3 loop of {
m HIV}\text{-1} gpl20 - used to treat {
m HIV} infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 39; DB 2; Length 8; 100.0%; Pred. No. 1.88+06; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV gp120 V3 loop; liposome composition; HIV infection.
                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 11; 39pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB68603 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV gp120 V3 loop peptide #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8; Conservative
                                                                                                                                                                                                                                                                                                    WPI; 1995-131312/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RAFVTIGK
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JS6180134-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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The present invention relates to a liposome composition comprising liposomes having an outer surface layer of polyethylene glycol chains, each having a free distal end. A polypeptide or polysaccharide effector molecule is covalently attached to a portion of the distal ends. The effector interferes with specific binding of pathogen or cell in a colorable composition may be used in treating a condition mediated by inding a pathogen or cell in the bloodstream, to a target cell or cell matrix, and is rapidly removed by renal clearance from the bloodstream, to a target cell or cell inding a pathogen or cell in the bloodstream, to a target cell or cell matrix. It can be used in treating septic shock, toxic shock, colonic inflammation, leukaemic cell proliferation, or HIV infection. The present sequence is a peptide of the V3 loop of HIV envelope protein gpl20. This compute to the CD4 receptor during HIV infection of lymphocytes. By introducing the present peptide, the CD4 receptors are blocked, thereby introducing HIV infection. (Updated on 11-SEP-2003 to standardise OS
                                                                                                                                                                          Liposome composition for use in treating septic shock comprises liposomes having an outer surface layer of polyethylene glycol chains, and a polypeptide or polysaccharide effector molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 39; DB 4; Length 8; 100.0%; Pred. No. 1.8e+06; ive 0; Mismatches 0; Indels
                                                                                                            Woodle MC, Martin FJ, Barenholz Y,
                                                                                                                                                                                                                                      Disclosure, Fig 13; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR46523 standard; peptide; 9 AA
95US-00480332.
                               93US-00035443
                                              94US-00316436
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(first entry)
                                                                            (SEQU-) SEQUUS PHARM INC
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Best Local Similarity
--han 8; Conserve
                                                                                                                                          WPI; 2001-201897/20.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 8 AA;
07-JUN-1995;
                               23-MAR-1993;
                                              29-SEP-1994;
                                                                                                            Zalipsky S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
30-MAR-1994
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The peptide is an example of an antigenic peptide which may be joined via a functional linker group, opt. at the two ends of a spacer group, to a functional linker group, opt. at the two ends of a spacer group, to a polar injection arrier, e.g. sphingosine, oeramide, phosphatidic coid. The compsn. may be used as a vaccine against HIV infection. An advantage of the campsn. may be used as a vaccine against HIV infection. An advantage of the arrier system is that when incorporated into the compsn. entry of the antigenic peptide into the cells of the immune system is facilitated (no need for endocytosis) and targetting to specific organelies becomes possible. Unlike known vaccines, intracellular synthesis of viral antigens is not necessary for presentation via the MHC class I antigen pathway, so both humoral and cellular immunity is achieved. I antigen pathway, so both humoral and cellular immunity is achieved. Also, when a spacer is present, the antigen release rate may be controlled. See also AAR46507-47. (Updated on 25-MAR-2003 to correct PF (field.) (Updated on 25-MAR-2003 to correct PF)
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                                                                                            New covalent conjugate of antigenic peptide and polar lipid e.g. sphingosine - useful in protective vaccines, treatment of auto-immune disease and preventing of transplant rejection.
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100.0%; Pred. No. 1.8e+06;
Live 0; Mismatches 0;
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                                                                                                                                                                                                                               Disclosure; Page 13; 15pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
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                            WPI; 1993-350862/44
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Stowell MHB, Malkovsky M;

Yatvin MB,

(OREG-) STATE OF OREGON.

92US-00911209 90US-00607982

09-JUL-1992; 01-NOV-1990; ប្រមូល ប្រមូល ò 용 ô immunological cytotoxic Tlymphocyte (CTL) response in a mammal. The method comprises: (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal's lymphatic system to maintain the immunologic CTL response. The method can be used for the delivery of e.g. a differentiation antigen, a tumour-specific multilneage antigen, an embryonic antigen, an oncogene antigen, a mutated tumour-suppressor gene antigen, or a viral antigen. They can be used for the treatment of disease such as cancer, e.g. malignant melanoma or infectious disease, e.g. viral disease such as hepatitis or AIDS. Sustained antigen delivery to the lymphatic system provides for potent CTL stimulation that takes place in the milieu of the lymphoid organ, and it sustains stimulation The UI snRNP is the target of high-titre, high avidity autoantibodies cocurring in the systemic rheumatooid disorders of mixed connective tissue disease, scleroderma and systemic lupus erythematosus. It has been found that some sites in the UI snRNP 70K protein (see AAR62120-R62135) are homologous to sites in HIV-1 gpl20/41 (AAR62136-R62125) and that anti-RNP autoantibodies can be used to neutralise HIV-1 (Updated on 25-MAR-2003 to correct DN fileld.) (Updated on 27-AUG-2003 to correct OS fileld.) The present invention describes a method of inducing and/or sustaining an Inducing a cytotoxic T lymphocyte response - by maintaining a level of antigen in the lymphatic system of a mammal so as to provide a sustained CTL response, used to treat, e.g. AIDS. Gaps Cytotoxic T-lymphocyte response; CTL; antigen; lymphatic system; immunisation; tumour; infectious disease; immunotherapy; cancer; malignant melanoma; viral disease; hepatitis; AIDS. ö ; DB 2; Length 9; 1.8e+06; 0; Indels 0; Mismatches 100.0%; Score 39; 100.0%; Pred. No. T cell epitope/MHC ligand SEQ ID NO:87. Disclosure; Page 54; 106pp; English. Disclosure; Page 26; 199pp; English. AAY10157 standard; peptide; 9 AA CTLI-) CTL IMMUNOTHERAPIES CORP. Human immunodeficiency virus 1. 98WO-US014289. 97CA-02209815 97US-00988320 12-MAY-1999 (first entry) 8; Conservative Kuendig TM, Simard JJL; WPI; 1999-120514/10. 1 RAFVTIGK 8 Best Local Similarity Sequence 9 AA; WO9902183-A2 10-JUL-1998; 10-DEC-1997; 10-JUL-1997; 21-JAN-1999 Synthetic. AAY10157; Query Match Matches ઠે

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The invention relates to methods and reagents for generating a protective CD8+ T-cell immune response against at least one target antigen. The kits CD8+ T-cell immune response against at least one target antigen. The kits core the invention comprises (1) as priming composition, a source of, one or more CD8+ T-cell [Cytotroxic T lymphocytes (CTL)] epitopes of the target antigen, plus a carrier and (41) as boosting composition a source of CTL epitopes, with at least one CTL epitope the same as used in (1), with this source being a non-replicating or replication-impaired recombinant poxvirus vector (PVV) plus a carrier. If the source of CTL epitopes in (1) is a viral vector, then the vector in (11) is from a different virus. The kits are used to generate an immune response (prophylactic or therapeutic) against pathogens or tumours, specifically against malaria parasites such as P. falciparum, or HIV, and also many other bacterial, craponse against melanome and cancer of breast or colon, and generally wherever a strong CD8+ response is protective. The boosting composition my be used alone to boost a naturally primed response against malaria. The specified PVV provide an excellent booster effect, better than that from wild-type poxvirus, resulting in complete rather than partial protection against epotozoite challenge. Also PVV are safer to use than
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that is necessary to keep CTL active, cytotoxic and recirculating through the body. AAY10071 to AAY10639 represent examples of peptide antigens given in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD8+ T-cell; immune response; antigen; priming composition; CTL; epitope; cytocoxic T lymphocyte; boosting; poxvirus vector; PVV; pathogen; tumour; malaria; parasite; P. falciparum; viral; bacterial; parasitic; cancer; melanoma; HIV; breast; colon; vaccination.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Generating CD8-positive T cell response to target antigen using recombinant poxvirus - for treating or preventing malaria and HIV infection, also epitope strings from Plasmodium and HIV.
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                                                                                                                                 Length 9;
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                                                                                                                               Query Match 100.0%; Score 39; DB.2; L
Best Local Similarity 100.0%; Pred. No. 1.8e+06;
Matches 8; Conservative 0; Mismatches 0;
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Hanke T, Smith GL, Blanchard T;
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                                                                                                                                                                                                                                                                                                                                                                            AAY03692 standard; peptide; 9 AA.
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07-JUN-1999
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Length 9; Indels

100.0%; Score 39; DB 5; I 100.0%; Pred. No. 1.8e+06; 0; Mismatches

Query Match Best Local Similarity 100. Matches 8; Conservative

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wild-type virus. Sequences AAY03681-704 represent CTL peptide epitopes of the HIV/SIV (H) epitope string. (Updated on 17-OCT-2003 to standardise OS
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                                                                       Gaps
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                                                      Length 9;
                                                                      0; Indels
                                                                                                                                                                                                              HIV epitope, HIV-1 gpl20 B*2705, peptide sequence.
                                                    ; Score 39; DB 2; I
; Pred. No. 1.8e+06;
0; Mismatches 0;
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                                                    100.0%;
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RAFVTIGK 9
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Best Local Similarity
Matches 8; Conserv
                                    Sequence 9 AA;
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AAU96033
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Major histocompatibility complex, MHC; MHC class I molecule, virus; epitope; cytocoxic T lymphocyte response; CTL response; lymphatic system; antigen; immunogenic; malignant tumour; carcinoma; melanoma; leukaemia; lymphoma; infectious disease; hepatitis; malaria; measles; tuberculosis; acquired immune deficiency syndrome; AIDS.

Human immunodeficiency virus.

WO200262368-A2

15-AUG-2002

MHC class I molecule, viral epitope #87.

15-NOV-2002 (first entry)

ABG79839;

ABG79839 standard; peptide; 9 AA.

RESULT 9 ABG79839

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The invention relates to a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal comprising administering directly to the lymphatic system of the mammal:

(a) an antigen in the form of a polypeptide; (b) a vector comprising a method is useful for inducing and/or sustaining CTL response in a mammal. This is particularly useful for treating a mammal having a malignant tumour (e.g. carcinoma, melanoma, leukaemia or lymphoma) or infectious disease (e.g. hepatifis, acquired immune deficiency syndrome (AIDS), malaria, measles or tuberculosis), or in an animal having a predisposition to these diseases. The mammal may be dogs, cats, mice, cattle, sheep, pigs, goats, rabbits, or preferably humans. ABG79753.

ABG80319 represent viral epitopes on major histocompatibility complex (MHC) class I molecules, used in the method of the invention
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The present invention relates to a new vaccine comprising a noncepticating vaccine delivery vehicle (which comprises a non-infectious
repolicating vaccine delivery vehicle (which comprises a non-infectious
recombinant viral tubule) carrying one or more immunogens. The invention
is useful for inducing an immune response, preferably danti-viral, in a
cubject. The administration of the vaccine is preferably followed by
caministering one or more virus like particles carrying an immunogen. It
is also useful for administering to a patient for generating antibodies
ce specific for one or more immunogens, such as Bluetongue virus (BTV), foot
and mouth disease virus (FMDV), influenza virus and human
immunodeficiency virus (HIV). The invention provides an effective means
of delivering multiple peptide components representing viral/tumour
contigenic groups to elicit protective immunity, which has not previously
contigenic groups to elicit protective immunity, which has not previously
contigenic groups to elicit protective immunity, which has not previously
contigenic groups to elicit protective immunity which has not previously
collection (AAU96022-AAU96045) of HIV epitopes that were used in the

methods of the invention as immunogens. These epitopes were used in construct chimeric NS1-TUBs (virus-derived tubules) which show immunogenicity. (Updated on 29-AUG-2003 to standardise OS field)

Sequence 9 AA;

Inducing or sustaining immunological cytotoxic T lymphocyte response in a mammal, useful for treating a mammal with malignant tumor or infectious disease, by directly administering an antigen to the lymphatic system of

(CTLI-) CTL IMMUNOTHERAPIES CORP. 02-FEB-2001; 2001US-00776232.

Simard JJL;

Kundig TM,

WPI; 2002-657506/70.

27-SEP-2000; 2000US-0235614P

(UABR-) UAB RES FOUND

WPI; 2002-339987/37.

Roy P;

22-JAN-2002; 2002WO-US002033

Disclosure; Page 20; 73pp; English.

A vaccine, for inducing an antiviral immune response, comprises a non-replicating vaccine delivery vehicle (which comprises a non-infectious recombinant viral tubule) carrying one or more immunogens.

Claim 8; Page 39; 65pp; English.

the mammal

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Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RAFVTIGK
                                                                                        WO2003104428-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2004071457-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9 AA;
                                             Unidentified
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                                                                                                                                 18-DEC-2003
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HLA-B*2705.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of generating an immune response against a non-hepadnaviral antigen in a mammal by administering (to the mammal) at least 1 dose of a priming component comprising a recombinant hepatitis B core particle (rHED) (which is a carrier for 1 or more non-hepatitis B core particle (rHED) (which is a carrier for 1 or more non-hepadnaviral CD8+ T-cell epitopes of the antigen). The method may be supplemented by the use of a boosting stage comprising a non-replicating or replication-impaired recombinant poxvirus vector. The method is useful for generating an immune response against a non-hepadnaviral antigen in a mammal for treating or preventing cancer or malaria. This sequence represents a peptide from human immunodeficiency virus type I (HIV-1) used as a carrier peptide to augment the immune response against a
                                                                                                                                                                                              HIV1 carrier peptide to augment anti-malaria CD8+ T-cell immune response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Generating an immune response against a non-hepadnaviral antigen in a mammal, useful for treating or preventing cancer or malaria, by administering a priming component comprising a recombinant hepatitis B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                         antimalarial; cytostatic; vaccine; immune response;
non-hepadnaviral antigen; hepatitis B core particle; CD8+ T-cell;
epitope; poxvirus vector; cancer; malaria; epitope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Breast/bladder carcinoma-related HLA-B*2705-binding peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 39; DB 7; Length 9; 100.0%; Pred. No. 1.8e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 49; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYNY-) UNIV NEW YORK MEDICAL CENT.
                                                                 ADE79993 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK50933 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-PEB-2003; 2003WO-US003897.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-FEB-2002; 2002US-0354963P
                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Birkett AJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-748124/70.
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                                                                                                                                                                                                                                                                                                                                                                           WO2003066833-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      core particle.
                                                                                                                                                     29-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                   14-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zavala F,
                                                                                                            ADE79993;
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                      ADE79993
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The invention relates to a novel isolated polypeptide comprising or consisting of two or more C35 peptide epitopes. The polypeptide of the invention demonstrates cytostatic activity and may be useful for the formulation of an immunogenic composition, such as a vaccine, to induce antibodies and cell-mediated immunity against target cells such as tumour cells. Furthermore, the polypeptide and its analogues may be useful as prognostic markers for carcinoma, such as human breast or bladder carcinoma. The current sequence is that of breast/bladder carcinomarelated HLA-B*2705-binding peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel C35 polypeptide useful for formulation of immunogenic composition to induce antibodies and cell-mediated immunity against tumor cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 12; Page 493; 626pp; English.
                                                                                                                                                                                                                                                                                                                                      Borrello MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADR69467 standard; peptide; 9 AA.
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18-APR-2003; 2003US-0463746P.
16-SEP-2003; 2003US-0503417P.
10-JUN-2003; 2003WO-US018252
                                                                            10-JUN-2002; 2002US-0386738P
                                                                                                           11-DEC-2002; 2002US-0432241P
23-APR-2003; 2003US-0464650P
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                                                                                                                                                                                                                                                                                                                                      Zauderer M, Evans EE,
                                                                                                                                                                                                                       (VACC-) VACCINEX INC. (UYRP ) UNIV ROCHESTER
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Claim 10; Page 119; 154pp; English.

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This invention relates to a novel hybrid antigen which comprises at least one antigenic domain of an infectious agent or tumour antigen and binding domain that non-covalently binds to a heat shock protein. The invention may be useful for the production of compounds with an antimicrobial or cytostatic activity. In addition, the invention may prove useful for the production of a vaccine or for gene therapy. The composition and methods disclosed are useful for preventing or treating infectious diseases or cancer. The present sequence is that of a HIA-A2 binding peptide which was used in the exemplification of the invention.
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                                                                                                                                  Slusarewicz P, Andjelic S;
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                                                                                                                                                                                                                                                                                         New hybrid antigens comprising an antigenic domain and improved behock protein-binding domains, useful for preventing or treating infectious diseases or cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 39; DB 8; Length 9; 100.0%; Pred. No. 1.8e+06;
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                                                                                                                           Fletchner J, Prince-Cohane K, Mehta S,
                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 115; 56pp; English.
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                                                                 (MOJA-) MOJAVE THERAPEUTICS INC.
12-FEB-2004; 2004US-00776521.
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nes 8; Conservative
                                                                                                                                                                                                                               WPI; 2004-625768/60.
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New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency virus infection.

(AMNA-) AMERICAN NAT RED CROSS.

Scott D, Zambidis E;

WPI; 1998-506315/43.

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                                             This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, 1gH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, as prevents or inhibits 'bystander' apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral T cell immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hydrophilic; antigenic determinant; HIV; envelope; glycoprotein; env; gp; recognition; B lymphocyte; type specific; antibody; vaccine; protection; immune response; infection; neutralisation; hypervariable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic human immunodeficiency virus vaccine - comprising hydrophilic peptide corresponding to at least 1 antigenic determinant of envelope glyco:protein recognised by B lymphocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 39; DB 2; Length 10; 100.0%; Pred. No. 0.086; o. Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1997-244862/22.
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                            Sequence 10 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-FEB-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW19056;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated N-acetyl-galactosaminyl-transferase enzyme - used for the production of glycosylated polypeptide(s) having particular enzymatic, immunogenic or other biological or physical properties.
                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                           polypeptide N-acetylgalactosaminyltransferase; GalNAc-t3; human; glycosylation; HIV gpl20.
composition comprising HIV env peptides that disrupt gp120/gp41
                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                     Score 39; DB 2; Length 11; Pred. No. 0.095; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                               UDP-N-acetyl-alpha-D-galactosamine;
                                                                                                                                                                                                                             AAW34472 standard; peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 4; 70pp; English
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                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus.
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                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                   8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Clausen H, Bennett EP;
                                                                                                                                                                                                                                                                                                                   gp120 V3 peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CLAU/) CLAUSEN H.
(BENN/) BENNETT E P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-008874/01.
                                                                                                                            1 RAFVTIGK 8
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8; Conserva
                                                                       Query Match
Best Local Similarity
                                           Sequence 11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 11 AA;
                interactions
                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9743405-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .S-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAY-1996;
                                                                                                                                                                                                                                                                                       11-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-NOV-1997.
                                                                                                                                                                                                                                                         AAW34472;
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Best Local &
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Matches
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ID AAW
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                                                                                                                                                                                                                                                                                                                                                                                                         autoantibody; immunoinfective cluster virus; nuclear protein antigen; systemic rheumatic disorder; human immunodeficiency virus; HIV-1; systemic lupus erythematosus; mixed connective tissue disease; scleroderma; glycoprotein 120; glycoprotein 41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Methods for treating immunoinfective cluster virus infections - utilis antibodies or fragments characteristic of auto antibodies produced by patients with rheumatic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                               epitope;
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                                                                                                                                                                                                                                                                                                                                                                           Small ribonucleoprotein complex; Ul snRNP; 70K protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV gp120 envelope protein, peptide 127, analogue 267d
                                                                                                                                                                                                                                                                                                                 HIV-1 gp120/41 protein consensus binding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Douvas A, Takehana Y, Ehresmann G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 56; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW54932 standard; peptide; 12 AA.
                                                             AAR62152 standard; peptide; 12 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYSC-) UNIV SOUTHERN CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus 1.
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                                                                                                                                                                                  (revised)
(revised)
(first entry)
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-MAR-1993;
                                                                                                                                                                                  27-AUG-2003
25-MAR-2003
02-MAY-1995
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                                                                                                                           AAR62152;
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AAR 62152
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AAR 6215
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for passive immunisation against infection.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Epitope; human immunodeficiency virus type 1; HIV-1; gp120; gp160; monoclonal antibody; V3 loop; immunisation; mouse; splenocyte; hybridoma; membrane fraction; passive immunisation; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monoclonal antibody specific for human immunodeficiency virus type 1 MN
                                                                                                                                                                                                                                                                                                                                                                            Gaps
Immunoadsorbent; immunoassay; HIV gp120; immunogen; antibody; Human.
                                                                                                                                                                                                                               HIV gp120 peptides - useful as immunoassay reagents or vaccine
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                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 39; DB 2; Length 12; 100.0%; Pred. No. 0.1; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV-1 strain IIIB gp120 V3 loop peptide.
                                                                                                                                                                                                                                                              Example 8; Column 21/22; 34pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW22327 standard; peptide; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus 1.
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                                                                                                         88US-00155321.
91US-00663262.
91US-00726605.
94US-00326676.
                                                                                                                                                              (UNBI-) UNITED BIOMEDICAL INC
                    Human immunodeficiency virus
                                                                                      95US-00488252
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Best Local Similarity 100.00
Best Local Similarity 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                1 RAFVTIGK
                                                                                                                                                                                                                                                                                                                               Sequence 12 AA;
                                                                                                                    01-MAR-1991;
09-JUL-1991;
19-OCT-1994;
                                                                                     07-JUN-1995;
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                                          US5763160-A.
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18-SEP-1997
                                                               09-JUN-1998
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                                                                                                                                                                                                                                             components.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW22327;
                                                                                                                                                                                    Wang CY;
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The invention relates to a novel monoclonal antibody (MAD) NMO3 which binds to epitopes from the human immunodeficiency virus type 1 (HTV-1). The antibody was raised conventionally by immunising Balb/C mice with purified live HTV-1 MM, then isolating splenocytes and fusing them to P3-K63-Age-U1 cells. Hybridomas were then screened with membrane fractions from infected and non-infected H9 cells. The MAD was observed to bind to a protein band of 120 kD on a Western blot of separated, denatured HIV-1 porteins. This binding was shown to be between residues 320-327 by epitope mapping by ELISA and competitive binding. The ability of the MAD to inhibit infection of cells by HIV-1 shown by infecting H9 cells with sequence represents the V3 loop region from HIV-1 strain in IIIS, where the MAD NMO3 binds. The MAD can be used for the passive immunisation of bumans susceptible to, or infected with HIV-1. (Updated on 17-OCT-2003 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monoclonal antibody which binds to HIV-1gp120 or gp160 - used to prevent and treat human HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW62889-900 represent peptides used to identify a peptide sequence (AAW62874) present in HIV-1gp120 or gp160 which is bound by the monoclonal antibody of the invention. The antibody neutralises in vitro the infection of H9 cell by an active HIV strain MN according to the p24 analytical method. The antibody is used for treatment of human HIV
                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monoclonal antibody, HIV-1gp120; gp160; infection; H9 cell;
HIV strain MN; treatment; human HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                        Score 39; DB 2; Length 13;
Pred. No. 0.11;
                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peptide seguence of the specification.
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                               Example 3; Col 10; 14pp; English
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Best Local Similarity 100.0%;
Matches 8; Conservative 0
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                                                                                                                                                                                                                                                                                                                  standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 RAFVTIGK
                                                                                                                                                                                                                                                                                                                                                       Sequence 13 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP10182489-A.
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100.0%; Score 39; DB 2;

Query Match

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DB 4; Length 13;

Score 39; DB 4; Pred. No. 0.11; 0; Mismatches

n 100.0%; Similarity 100.0%; 8; Conservative 0

Query Match Best Local S Matches 8

RAFVTIGK 13

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8

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stabilized pharmaceutical containing N-terminal glutamic acid or glutamine, useful e.g. in anti-melanoma vaccines, is an addition salt
                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention describes a pharmaceutical compound (I) that
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                                           0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vaccine related MHC ligand peptide SEQ ID NO:536.
            Pred. No. 0.11;
100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Klinguer-Hamour C, Corvaia N, Beck A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (FABR ) FABRE MEDICAMENT SA PIERRE.
                                                                                                                                                                                                                                                                                                             AAM99433 standard; peptide; 13 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 9; Page 122; 149pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-MAR-2001; 2001WO-FR000872.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-MAR-2000; 2000FR-00003711
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                           8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
                                                                                                                                       |||||||||
RAFVTIGK 13
                                                                                                     1 RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-611470/70
            Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    glutamine, useful
with strong acid.
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07-DEC-2001
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Sequence of peptide which corresp. to the V3 loop region of gpl20 of HIV-1 isolate IIIB.

(revised)
(first entry)

25-MAR-2003 06-JUL-1993

AAR33336;

AAR33336 standard; peptide; 14 AA.

RESULT 21 AAR33336 ID AAR3 Monoclonal antibody, NM-01; HIV-1; gp120; gp160.

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MN-01 is a monoclonal antibody. In order to characterize the viral epitope recognized by NM-01, the antibody was screened by ELISA for exactivity with overlapping peptides corresp. to the amino acid sequence of the V3 loop region of HIV-1 gpl20 (AAR33332, AAR3333, AAR33334).

While there was no detectable reactivity over background of MAb-01 with the peptides corresp. to AAS 302-316 or 132-336 of the V3 loop, binding of the antibody to the peptide representing AAS 312-326 was apparent. The extent of this reactivity with other HIV-1 isolates was screened with peptides corresp. to the V3 loop region of HIV-1 isolates IIIB, RF, CDC4, NY/5, Z6, Z2 and ELI (AAR33335-R33342). These results indicate that monoclonal antibody NM-01 recognizes an epitope of the V3 loop of gpl20 of multiple HIV-1 isolates having the amino acid sequence AAR33334.

The variable region of the heavy and light chain of monoclonal antibody NM-01 were cloned by PCR and sequenced. Nucleotides 1-21 and 334-363 of AAQ3742 corresp. to the PCR primers used to amplify NM-01 light chain sequences and nucleotides 1-27 and 385-402 of AAQ57471 corresp. to the CDC4 MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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100.0%; Score 39; DB 2; Length 14;
Best Local Similarity 100.0%; Pred. No. 0.12;
Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 14 AA;
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contains an N-terminal glutamic acid (Glu) or glutamine (Glu) residue in the form of an addition salt with a strong, physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition containing at least one (I); (a) a containing at least one (I); (b) a containing at least one (I); (c) a method for in vitro diagnosis of diseases associated with the presence of (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process for preparing (I). (I) has immunomodulator, endocrine, antiallergic, neuroprotectant, virucidal, bactericidal, antiparasitic, fungicidal and cytostatic activities. (I) are useful, in human or veterinary medicine, in pharmaceutical compositions (for treating immune disorders, e.g.

immune deficiency, autoimmune states, hypersensitivity, allergy, graft rejection, infection, hormonal disorders and central nervous system diseases), also, where (1) is a MHC ligand (1a), in vaccines for treatment to prevention of: (1) viral, bacterial, parasitic or fungal infections; or (1i) of cancers. A particular application is in antimelanoma vaccines. (1) are also useful for in viro diagnosis of diseases associated with interactions between MHC and (1), e.g. melanoma and human immunodeficiency virus infection. AAM98898 to AAM99592 represent peptides which can be used in pharmaceutical compounds from the present invention. (Updated on 11-SBP-2003 to standardise OS field)

Sequence 13 AA;

Monoclonal antibodies against HIV-1 gp120 and gp160 proteins - for treating and preventing HIV-1 infection.

Example; Page 20; 57pp; English

NISP ) NISSIN SHOKUHIN KAISHA LTD

22-AUG-1991;

WPI; 1993-093943/11.

Ohno T,

92WO-US007111 91US-00748562

24-AUG-1992;

WO9304090-A1

Synthetic.

04-MAR-1993

1 RAFVTIGK ð

7 RAFVTIGK 14

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AAR48604 standard; peptide; 14 AA. (revised)
(first entry) 25-MAR-2003 03-SEP-1994 AAR48604; RESULT 22 

Human immunodeficiency virus; HIV-1; AIDS; glycoprotein; V3 loop; gp120; Sequence of portion of gp120 V3 loop peptide from HIV-1 isolate IIIB. epitope; isolate IIIB.

Human immunodeficiency virus 1.

WO9404574-A1

03-MAR-1994.

24-AUG-1993;

92WO-US007111. 24-AUG-1992; 22-APR-1993;

(NISP ) NISSIN SHOKUHIN KAISHA LTD.

Ohno T;

WPI; 1994-083117/10.

New humanised antibody specific for epitope on HIV-1 gp 120 - able to neutralise infection of HG cells, also nucleic acid encoding it, useful for passive immunisation to treat or prevent HIV-1 infection.

Example; Table 4, Page 18; 91pp; English.

GPGR is a portion of HIV-1 gpl20 or gpl60 protein. Monoclonal antibodies (MAbb) that react with this and which have the capacity to neutralise the infection of H9 cells in culture by live HIV-1 strains MN and IIIB are claimed. Specifically illustrating the invention are the murine MAD (designated NM-01) produced by hybridoma cell line HB 10726 which is deposited under ATCC NO. HB 10726, and the humanised versions of AD NM-01. To identify the specific epitope of gpl20 recognised by NM-01, the AD was screened for reactivity with three overlapping peptides corresp. to the V3 loop region of gpl20 (AAR4860-02). While there was no detectable reactivity over background of MAD NM-01 with the peptides corresp. to AAB 302-36 of the V3 loop, binding of the AD to the peptide creativity over background of MAD NM-01 with the peptides corresp. to AAB 302-36 of the V3 loop region of HIV-1NM 9120 prompted further studies on the extent of this reactivity with other HIV-1 isolates. The Ab was screened by BLISA for reactivity with peptides corresp. to the AB sequences of the peptides are given in AAR48603-10. NM-01 reacted with the loop peptides from the NM, IIB, RF, and CDC4 isolates. It showed a lesser affinity for the NV/5 peptide. (Updated on 25-MAR-2003 to correct DEST CONSTANT CONSTA

Sequence 14 AA;

Gaps . . 100.0%; Score 39; DB 2; Length 14; 100.0%; Pred. No. 0.12; ive 0; Mismatches 0; Indels Local Similarity 100. Query Match Matches

1 RAFVTIGK 8

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||||||||| ||RAFVTIGK 14

AAR66416 standard; peptide; 14 AA. 

AAR66416;

(revised)
(first entry) 25-MAR-2003 03-AUG-1995

HIV-1 IIIB peptide 18-1 (316-330).

human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.

Synthetic.

WO9426785-A1.

24-NOV-1994.

94WO-US005142. 13-MAY-1994;

93US-00060988 14-MAY-1993;

(USSH ) US SEC DEPT HEALTH.

Nara P, Ahlers JD, Pendleton CD, Berzofsky JA,

WPI; 1995-006707/01.

Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.

Example 1; Page 33; 120pp; English.

in the HIV-I IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR6416-R66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-1, the N-terminal residue (Arg) in peptide 18 has been deleted. (Updated on 25-MAR-2003 to correct Single residues from the HIV-1 RF sequence (AAR66415) replaced residues PN field.

Sequence 14 AA;

ö 100.0%; Score 39; DB 2; Length 14; 100.0%; Pred. No. 0.12; ive 0; Mismatches 0; Indels 8; Conservative Query Match Best Local Similarity Best Loc Matches

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Gaps

RAFVTIGK 14 1 RAFVTIGK 8

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RESULT 24

AAR66417 standard; peptide; 14 AA. 

AAR66417;

(first entry) (revised) 25-MAR-2003 03-AUG-1995

HIV-1 IIIB peptide 18-2.

T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20;

RESULT 23

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(NISP ) NISSIN SHOKUHIN KAISHA LTD
93US-00039457.
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                                                                                                                                                                                                                                                                                                                                                      7 RAFVTIGK 14
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                                                            WPI; 1996-442363/44
                                                                                                                                                                                                                                                                                                                                         1 RAFVTICK
                                                                                                                                                                                                                                                                               Sequence 14 AA;
22-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus sp.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9836087-A1.
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                                                                                                                                                                                                                                                                                                  Query Match
                                       Ohno T;
                                                                                                                                                                                                                                                                                                             Best Loc
Matches
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                                                                                                                                                                                                                                         Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66410, showed that binding was enhanced over peptide 18 control when a tyrosine was substant for a Val at position 11 and substants. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substans. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-2, the 11e residue at position 2 in peptide 18 has been deleted. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus type-1; HIV-1; gp120; gp160; epitope; monoclonal antibody; infection; heavy chain; light chain; hybridoma; complementarity determining region; CDR; V3 loop.
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                                                                                                                                           Shirai M;
                                                                                                                                                                                                                                                                                                                                                                                                       DB 2; Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 39; DB 2; Length 14 100.0%; Pred. No. 0.12; ive 0; Mismatches 0; Indels
cluster peptide; principal neutralising determinant.
                                                                                                                                           Berzofsky JA, Ahlers JD, Pendleton CD, Nara P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV-1 strain IIIB gp120 V3 loop peptide.
                                                                                                                                                                                                                            Example 1; Page 33; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW09264 standard; peptide; 14 AA
                                                                                94WO-US005142.
                                                                                                    93US-00060988.
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92WO-US007111.
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                                                                                                                       (USSH ) US SEC DEPT HEALTH
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(first entry)
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                                                                                                                                                              WPI; 1995-006707/01.
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                                                                                                                                                                                                                                                                                                                                                                                    Sequence 14 AA;
                                        WO9426785-A1
                                                                                13-MAY-1994;
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24-AUG-1992;
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03-MAR-1997
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                                                            24-NOV-1994
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                    Synthetic
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AAW09264
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                                                                                                                                                                                                                      The invention relates to a novel monoclonal antibody designated NM-01. The antibody was raised by immunising 2-month old Balb/c mice with live HIV-1 strain MN. Splenocytes from the mice were fused to P3-X63-Ag8-U1 cells (ATCC CRL1597). Hybridomas were screened using membranes from non-infected and MN-infected H9 cells, by reacting with hybridoma culture supernatants. This screening was followed by immunoflucrescence and radiodmmunoassays. The screening isolated the hybridoma HB 10726 which secretes the antibody NM-01. The peptides AAW09263-72 are derived from other HIV strains and were used to determine which other HIV-1 isolates antibody NM-01 reacted with. This peptide is from HIV-1 strain IIIB. The antibody is used for the diagnosis of HIV-1 in a fluid e.g. blood, and can be used to treat or prevent an HIV-1 infection. (Updated on 25-MAR-
\mu monoclonal antibodies to HIV-1 - used for the prevention, treatment diagnosis of HIV-1 infection.
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                                                                                                                                                    Example 2; Col 11; 56pp; English
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induce tolerance to the epitopes for treatment of human immune deficiency

virus infection

This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IGH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immunoglobulin heavy chains (HIV) gpilo epitopes. Fusion IG proteins and/or transfected cells are used to tolerise subjects to gpilo epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as proteins can be used against other diseases where an immune response is cleleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpilo, so prevents or inhibits 'bystander' apoptosis of uninfected T cells that are bound to gpilo protein, maximising induction of protective antiviral This peptide elicits cytotoxicity by T-cells against HIV antigens and stimulates prodn. of HIV-specific cytotoxic T-lymphocytes (CTLs). It is specific for the HIV envelope protein gp160. (Note: Revised entry submitted to correct the patent number formst of US Government-owned NTIS applications to prevent clashes with ongoing US granted patent numbers. Synthetic peptide corresp. to HIV GP 160 ENV sequence - which elicits cytotoxicity by T cells against HIV and proliferation of HIV-specific cytotoxic T cells. 100.0%; Score 39; DB 2; Length 14; 100.0%; Pred. No. 0.12; Germain R, Moss 0; Indels Env-K1; gp160 Env protein; T-cell cytotoxicity; HIV. Mismatches Berzofsky J, Takahashi H, Hosmalin A, (USSH ) US DEPT HEALTH & HUMAN SERVICE. (USDC ) US SEC OF COMMERCE. Claim 10; Page 119; 154pp; English AAP82095 standard; peptide; 15 AA. Disclosure, Page ?; 31pp; English. ; 0 88US-00148692. 88US-00148692. (revised)
(revised)
(first entry) 8; Conservative 5 RAFVTICK 12 1 RAFVTIGK 8 WPI; 1988-264280/37. Query Match Best Local Similarity Matches 8; Conserv Sequence 14 AA; Env-K1 peptide. 25-MAR-2003 17-DEC-2001 29-OCT-1990 USN7148692-N. 26-JAN-1988; 26-JAN-1988; 02-AUG-1988. Synthetic. AAP82095; RESULT 27 AAP82095 ò 요 

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For further information please visit the Derwent web site at www.derwent.com/dwpi/updates/ntis us.html.) (Updated on 25-MAR-2003 to correct PF field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic human immuno-deficiency virus env-coded peptide(s) - induce antibodies that block human immuno-deficiency virus proliferation and fusion between infected and non-infected cells.
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                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                      Peptide comprising AAs 308-322 of HIV-I IIIB env protein.
                                                                            100.0%; Score 39; DB 1;
100.0%; Pred. No. 0.13;
ive 0; Mismatches 0.
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(DUPO ) DU PONT MERCK PHARMACEUTICAL CO.
                                                                                                                                                                                                                     AAP91228 standard; peptide; 15 AA.
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89US-00324027.
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Best Local Similarity 100.
Matches 8; Conservative
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8 RAFVTIGK 15
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es 8; Conserv
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                                                   Sequence 15 AA;
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25-MAR-2003
13-AUG-1990
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Gaps ö RESULT 29 AAR06294

Chang NT;

Chang TW,

Fung MS,

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The inventors claim a chimeric, viral-neutralising immunoglobulin which binds to the gp120 region of HIV-1 with a potency and immunologic binds to the gp120 region of HIV-1 with a potency and immunologic antigen-binding region of non-human origian and a constant region of cantigen-binding region of non-human origian and a constant region of thuman origin. Specifically claimed is the chimeric immunoglobulin CGP 47439. Probes V-kappa-1 and V-kappa-2 (AA021497, AA021498) were used to screen a genomic DNA library for BAT123 sells for the functionally rearranged variable region en of the mouse/human chimeric L chain gene. Probe VH-1 was used to screen partial genomic libraries for the functionally rearranged variable region cof the mouse/human chimeric L chain gene. Probe VH-1 was used to screen partial genomic libraries for the functionally rearranged variable region coffice or this clone was used in the construction of the mouse-human confidence (AAR21343) as BAT123 which indicates that the conversion libraries (AAR21343) as BAT123 which indicates that the antigen specificity of the murine antibody BAT123 was preserved upon conversion libraries antigen specificity of the murine antibody (Updated on 25-MAR-2003)
                                                                                               New chimeric HIV-1-neutralising immunoglobulin(8) - comprising non-human antigen binding regions and constant human region, for immuno-therapy of AIDS and ARC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gp120; HIV-1; cytotoxic T-lymphocyte; CTL; T-helper; AIDS; infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.13;
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                                                                                                                                                                       Example, Page 26; 39pp; English.
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                                             Sun BN,
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                (TANO-) TANOX BIOSYST INC.
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Best Local Similarity 100.
Matches 8; Conservative
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                                                                         WPI; 1992-064897/08
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16-SEP-1992;
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                                           Liov RS,
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AAR38187
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BAT123 mouse immunoglobulin; viral antigen-binding region; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide is a fragment of HIV-1 gp 120 glyco-protein, useful in providing vaccines, preferably operatively linked to an immunogenic carrier eg. keyhole limpet haemocyanin, HSA, tetanus toxoid etc. (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1 gp120 epitope found in mouse immunoglobulin BAT123 and mouse/human chimeric antibody CAG1-51-4.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide(s) derived from HIV-1 - stimulate T-cell proliferation, etc.
useful in immunisation against HIV.
                                                                                                                                      AIDS; vaccine; T-cell proliferation; keyhole limpet haemocyanin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.13; ive 0; Mismatches 0; Indels
                                                                                                         Peptide derived from HIV-1 gp 120 envelope glycoprotein.
AAR06294 standard; protein; 15 AA
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                                                                                                                                                                     Human immunodeficiency virus 1.
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RAFVTIGK 9
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17-DEC-1990
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16-MAY-1992
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                              AAR06294;
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HIV gpl20 V3 loop-derived peptides (AAR38170-87) are successful in generating CTL responses, esp. peptide RISK (AAR38187); the T-helper cell-inducing peptide includes the sequence C19A (AAR38164); HIV infection-inhibiting peptides are given in AAR38188-206, and are esp. peptides RISK, N24G, E13V, R8K, T13Q and H13N (AAR38165-69). The peptides may also be derived from an influenza virus protein or a sendai virus protein (AAR41014-15). It was observed that peptide RISK (amino acids 315-329), with sequences derived from the V3 loop of HIV-1 IIIB, inhibits HIV-1 infection of primary human T cells by 92% at 1 microg/ml (ca. 0.4-0.6 microM). (Updated on 25-WAR-2003 to correct PN field.) (Updated on 27-AUG-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      V3 loop, envelope glycoprotein; gp160; HIV-1; prophylaxis; immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The peptide corresponds to amino acid residues numbered 315-329 in the V3 loop of the envelope glycoprotein gp160 of human immunodeficiency virus (HIV-1), as numbered by Ratner in the strain MN. It is useful for the prophylaxis or immunoteherapy of HIV-1 infection. It elicits an immunised subject ctytoxic T lymphocyte (CTL) activity against the corresp. clinical isolate of HIV-1. (Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent
                 position for treating and preventing viral infections - \rm L\textsc{-}lnducing epitope and HIV infection-inhibiting sequence or T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide(s) corresponding to the V3 loop of gp=160 of HIV-1 - elicit cytotoxic T lymphocyte(s) active against broad range of HIV-1 isolate(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    quence of peptide which corresp.to AA residues 315-329 of the V3 loop the gp160 envelope glycoprotein in HIV-1 strain MN.
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                                                                                         Claim 13 + 19; Page 94-95; 130pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide which corresp.to AA
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                 Peptide composition for treating comprise CTL-inducing epitope and
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                                     comprise CTL-inducing epitope helper cell-inducing sequence.
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RAFVTIGK 15
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Matches 8; Conserv
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clashes with ongoing US granted patent numbers. For further information please visit the Derwent web site at www.derwent.com/dwpi/updates/ntis_us.html.) (Updated on 24-OCT-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                  120; HIV epitope; Human Immunodeficiency Virus fusion polypeptide.
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Pred. No. 0.13;
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Similarity 100.0%;
8; Conservative 0;
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(revised)
                                                                                                                                                                           RAFVTIGK 15
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Best Local Similarity
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                                                                                                                                                   1 RAFVTIGK
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                                                                    Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAR-1994.
                                                                                                                                                                                                                                                                                                    27-AUG-2003
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Best Local S:
Matches 8;
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                                                                                                                                                                                                                   RESULT 33
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AAR74603
ID AAR74
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Gaps

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AAR74603;

Shirai M;

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Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAP). In peptide 18-5, the Gly residue at position 5 in peptide 18 has been replaced by an Ala residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                             Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.13;
                                                                                                                                                                   Nara P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3erzofsky JA, Ahlers JD, Pendleton CD, Nara P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                   Pendleton CD,
                                                                                                                                                                                                                                                                                                                         Example 1; Page 33; 120pp; English.
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                                                     94WO-US005142
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                                                                                           93US-00060988
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                                                                                                                                                                   Ahlers JD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV-1 IIIB peptide 18-6.
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Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 AA;
                                                                                                                             (USSH ) US SEC
                                                                                                                                                                   Berzofsky JA,
                                                     13-MAY-1994;
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03-AUG-1995
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                 24-NOV-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MAD 5023 was developed against AA residue 308-322 of the variable loop of HIV-I (AAR74603). MAD 5023 binds to the epitope GRAF. G preceding RAF is believed to critical for internalization. MAD 5023 recognised p160, p120, p42 and p24 in cancer cells. AAR74603 competitively blocked binding of the MAD to the cancer antigens, indicating that at least the epitope GRAF, which is recognised by the MAD, must also be present in cancer antigens. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PI field.) (Updated on 16-OCT-2003 to standardise OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detection of HIV-1 cross-reactive breast carcinoma-associated antigens - for diagnosis and anti:sense therapy of breast and gynaecological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T cell helper site, cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                 MAb 5023; variable V3 loop; HIV-I; human immunodeficiency virus; cancer antigen; monoclonal antibody.
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                                                                                                                             HIV-I variable loop residues 308-322.
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                                                                                                                                                                                                                           Human immunodeficiency virus; I.
                                                                                                                                                                                                                                                                                                                                                                                                                       (RAKO/) RAKOWICZSZULCZYNSKA E M.
                                                                                                                                                                                                                                                                                                                                             94WO-US011754
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                                                                                           (first entry)
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(first entry)
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B; Conservative
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                                                     (revised)
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Shirai M;

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Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
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Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.

Example 1; Page 33; 120pp; English

Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-tyr66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of determinant sequence (PGRAF). In peptide 18-6, the Pro residue at position 6 in peptide 18 has been replaced by an Ala residue. (Updated on 25-MAR-2003 to correct PN field.)

Sequence 15 AA;

100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.13; , 0.13; has 0; Indels Mismatches ; 0 Conservative 8 RAFVTIGK 15 œ Local Similarity les 8; Conserv 1 RAFVTIGK

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Gaps ô

AAR66414 standard; peptide; 15 AA. (revised)
(first entry) 25-MAR-2003 AAR66414;

HIV-1 IIIB peptide 18.

T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant; IIIB isolate.

Synthetic.

94WO-US005142. 13-MAY-1994; 93US-00060988 14-MAY-1993;

(USSH ) US SEC DEPT HEALTH.

Shirai M; Nara P, Berzofsky JA, Ahlers JD, Pendleton CD, WPI; 1995-006707/01.

Example 1; Page 33; 120pp; English.

Synthetic peptides spanning multideterminant regions from the HIV envelope protein gp160 have been designed and are designated cluster peptides (PCLUS). These peptides each consist of a cluster of overlapping determinants and are known to induce in vitro T cell proliferation and cytokine production in mice and humans of multiple MHC types. The cluster

part of HIV ö T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant. Gaps immunodominant CTL determinant, Peptide 18 (AAR66414), corresp. to of the gpl60 V3 loop and principal neutralising determinant region -1 IIIB isolate. (Updated on 25-MAR-2003 to correct PN field.) ö Score 39; DB 2; Length 15; Pred. No. 0.13; 0; Indels the N-terminus 0; Mismatches peptides were co-linearly synthesised at AAR66419 standard; peptide; 15 AA. 100.0%; 94WO-US005142. 93US-0006098B. (USSH ) US SEC DEPT HEALTH. (revised)
(first entry) 8; Conservative HIV-1 IIIB peptide 18-4. 15 8 Local Similarity 1 RAFVTIGK 8 RAFVTICK Sequence 15 AA WO9426785-A1 13-MAY-1994; 14-MAY-1993; 25-MAR-2003 03-AUG-1995 24-NOV-1994. Synthetic. AAR66419; Query Match Matches RESULT 38 8\$88888 셤 à

Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66430) and AAR66411). Analysis of the substituted peptides (AAR66416-R66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of determinant sequence (PGRAF). In peptide 18-4, the Arg residue at position 4, in peptide 18 has been replaced by a Lys residue. (Updated on Gaps 25-MAR-2003 to correct PN field.) Sequence 15 AA;

Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.

Example 1; Page 33; 120pp; English.

Σ

Shirai

Nara P,

Berzofsky JA, Ahlers JD, Pendleton CD,

WPI; 1995-006707/01.

ö Query Match
100.0%; Score 39; DB 2; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.13;
Matches 8; Conservative 0; Mismatches 0; Indels

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1 RAFVTIGK 8

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AAR68744-805 are cytotoxic T lymphocyte (CTL) class I and II restricted epitopes derived from human immunodeficiency virus proteins. AAR68789 corresponds to amino acid residues 308-32 of the env gpl20 protein. These antigens are examples of peptides that can be used with an immunogenic implant. The implant is associated with an antigen associated with a pathogen and used to form a discrete, localised chronic inflammation site which acts as a local 'factory' for prodn. Of CTL's which lyse cells infected with a specific pathogen. The expanded set of pathogen-specific CTL's can eradicate or prevent development of infection, and can also be used to treat or arrest the development of cancers associated with infection. (Updated on 25-WAR-2003 to correct PN
                                                          cytotoxic T lymphocyte; epitope; antigen; pathogenic; nef; gag; pol; env; gpl20; gp41; HIV; cell-mediated immunity; human immunodeficiency virus; class I restricted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Implant for sustained release of pathogen-associated antigen - forming chronic inflammatory site producing cytotoxic T-lymphocytes lysing infected cells, esp. for treating AIDS.
                     Cytotoxic T lymphocyte epitope 46 derived from env gp120 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 100.0%; Score 39; DB 2; Length 15; Similarity 100.0%; Pred..No. 0.13; 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus 1; strain HXB2R.
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                                                                                                                                              Human immunodeficiency virus.
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(first entry)
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17-JAN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-AX66409) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAP). In peptide 18-7, the Gly residue at position 7 in peptide 18 has been replaced by an Ala residue. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                             T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gpl20; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Berzofsky JA, Ahlers JD, Pendleton CD, Nara P, Shirai M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR68789 standard; peptide; 15 AA.
                                                                                                                          AAR66422 standard; peptide; 15 AA
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(first entry)
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(first entry)
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                                                                                                                                                                                                                                                                         HIV-1 IIIB peptide 18-7.
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RAFVTIGK 15
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Matches 8, Conserv
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                                                                                                                                                                                                          25-MAR-2003
03-AUG-1995
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23-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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RESULT 40

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Gaps

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                                                                                                                                                                                   gClq receptor-based, HIV-1 gp 120 binding peptide(s) - for preventing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Epimorphin; human, mouse, wound; burn; epithelial tissue, diagnosis, treatment; morphogenetic abnormality; cosmetic; hair growth stimulator.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 15;
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100.0%; Pred. No. 0.13;
ive 0; Mismatches 0; Indels
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                                                                                              Kim YW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR92033 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                           Claim 10; Page 49; 53pp; English.
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95JP-00099979.
95JP-00099980.
                                                                                              Sun CRY,
                                                                                                                                                                                                              and treating HIV-1 infection.
95US-00410360
                                            (TANO-) TANOX BIOSYSTEMS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95EP-00304270
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Best Local Similarity 100.
Matches 8; Conservative
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                                                                                            Sun BNV,
                                                                                                                                        WPI; 1996-455274/45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 RAFVTIGK
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24-MAR-1995;
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31-MAR-1995;
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                                                                                            Fung MSC,
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New polypeptides contain a first portion of 5-99 amino acids joined to a second portion contg. at least a functional domain of epimorphin. The

Claim 8; Page 57; 62pp; English.

artificial organs.

Novel polypeptide containing an epimorphin functional domain - has possible benefits in epithelial tissue treatments, e.g. burns and for

WPI; 1996-118213/13.

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first portion may be selected from the peptides given in AAR92029 to AAR92036. The second portion may be full-length epimorphin (see AAR92037 to AAR92042 for human and mouse epimorphins). Fragments of epimorphins given in AAT16083 to AAT16090 are used in the prodn. of modified epimorphins are useful for the development of diagnosis and treatment of morphogenetic abnormalities of epithelial tissue or novel remedies for wounds, eg burns, after surgery and for artificial organs. They may also be used as components of cosmetics, hair growth stimulators, etc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence represents a HIV gpl20 peptide that can be used in the method of the invention. The method of the invention is for determining whether or not a mother will transmit HIV.1 to a feetus. The method comprises incubating asmple from the HIV-infected mother, with a collection of HIV peptides. The HIV peptides at least one for the gpl20 sequences (such as AAW07918-W07917), and at least one HIV gpt11 derived peptide (see AAW07918-W07928). The number of peptides that react with the sample is determined, and this number is compared with a standard that shows pattern reactivity for a patient of transmission status. A non-transmissive HIV sample is indicated if the test sample tacts with twice as many peptides as the standard. The method detects the presence of neutralising antibodies that protect against mother to infant transmission of HIV. These sequences can also be used in vaccines to protect against transmission. Antibodies against these sequences can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Determining the likelihood of maternal transmission of HIV-1 to foetus by measuring maternal reactivity with specific gp120 and gp41 derived peptide(s), also used for diagnosing HIV in infants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV; gpl20; HIV-IIIB strain; HIV-1 transmission; foetal transmission; neutralising antibody; passive immunisation; anti-idiotypic antibody; gp41; vaccine; active immunotherapy.
                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                              Similarity 100.0%; Score 39; DB 2; Length 15; Similarity 100.0%; Pred. No. 0.13; 8; Conservative 0; Mismatches 0; Thalo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gp120 peptide p18p.
                                                                                                                                                                                                                                                                                                                     1 RAFVTIGK
                                                                                                                                                                                                 Sequence 15 AA;
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31-JAN-1997
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AAW24219 standard; peptide; 15 AA

RESULT 45 AAW24219

AAW24219;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antigenic peptides can be inserted into the alpha-antigen sequence of a Mycobacterium and secreted from an appropriately transformed M.bovis BCG cell. The resulting chimeric antigen has greatly enhanced antigenicity and immunogenicity and is recognised in vivo by B-cells which recognise the alpha-antigen. The present sequence is that of a HIV-1 gpl20 V3 loop eptrope which was incorporated into the alpha antigen. M. bovis BCG cells secreting a chimeric protein comprising the epitope sequence are useful as anti-AIDS vaccines. (Updated on 16-OCT-2003 to standardise OS field)
be used for passive immunisation, and to generate anti-idiotypic antibodies for use in vaccines or active immunotherapy. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                     Mycobacterium bovis BCG; AIDS vaccine; surface protein; alpha antigen;
Human immunodeficiency virus type 1; fusion protein; gpl20 epitope;
                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                        HIV-1 V3 loop epitope, for insertion in Mycobacterium alpha antigen.
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                                                                                           DB 2; Length 15;
                                                                                                                       0; Indels
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                                                                                           100.0%; Score 39; DB 2;
100.0%; Pred. No. 0.13;
                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (AJIN ) AJINOMOTO CO INC.
(NINA-) JAPAN AGENCY NAT INST HEALTH.
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                                                                                                                                                                                                                                                              AAR92007 standard; protein; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
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Best Local Similarity 100.
                                                                                                                        8; Conservative
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                                                                                                                                                     1 RAFVTIGK 8
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                                                                                                        Local Similarity
                                                             Sequence 15 AA;
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27-SEP-1996
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Matches
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tracing sources of infectious diseases. The method comprises preparing a tracing sources of infectious diseases. The method comprises preparing a short (9-50 amino acid) peptide containing at least one non-conserved epitope of an organism, injecting a composition containing the peptide intradermally into a test subject in a delayed-type hypersensitivity (DTH) assay and observing the injection site at intervals for induration. In this example CD4+ Tlymphocyte epitopes to the HIV-1 V3 loop were mapped by existing in vitro technique for two existing HIV infected individuals and used to design peptides AAW24217-20. The method allows inclinately and used to design peptides AAW24217-20. The method allows and treatment design for infectious diseases, allergic diseases and malignancy. It is especially useful for tracing infectious diseases and malignancy. It is especially useful for tracing infectious diseases and malignancy. It is especially useful for tracing infectious diseases and malignancy in vivo mapping suitable vaccines. Vaccinated individuals can also be tested to verify protection against a particular strain. The method allows in vivo mapping of T-1 ymphocyte epitopes, not previously possible. The method is simpler, more rapid and more sensitive. It can also be applied in a variety of environments e.g. undevelopes since specialist equipment is not
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide fragments containing antigen epitope(s) used to trace diseases used in a delayed-type hypersensitivity assay, for in vivo mapping of human T-lymphocyte epitope(s) e.g. for diagnosis, vaccine development
                                                                                                CD4+ T-lymphocyte epitope to HIV-1 V3 loop derived peptide V3-LAI-P18.
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                                                                                                                                    T-lymphocyte epitope, diagnosis, antigen; infectious disease, delayed-type hypersensitivity assay; vaccine development.
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                                                                                                                                                                                                                     Human immunodeficiency virus.
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                                                        (first entry)
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8 RAFVTIGK 15
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                             27-JAN-1997;
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                                                        17-MAR-1998
                                                                                                                                                                                                   Synthetic.
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RESULT 46 AAW10348

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0; Indels

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RAFVTIGK 8

AAW10348;

Synthetic.

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96WO-US020463.
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                                                                                                                                                                                       (HARD ) HARVARD COLLEGE
                                                                                                                                                                                                                                                                             WPI; 1997-350782/32.
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                       Collier RJ,
Starnbach MN;
                            WO9723236-A1
                                                                                                 13-DEC-1996;
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IIIB for use in a method for diagnosing exposure of a patient to human
immunodeficiency virus (HIV). The method involves: (a) obtaining
peripheral blood mononuclear cells (BMC) from a patient; (b) incubating
the PBMC with at least 1 synthetic peptide representing an epitope(s) of
activation of the PBMC as a result of the incubation in step (b). The
method can provide for the early detection of exposure to infectious
organisms, specifically HIV in this case. The method can be used to
assess exposure to HIV without concomitant infection. It also provides an
earlier identification of HIV exposure, than is provided by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diagnosis of exposure to infectious agents, partic. HIV - by detecting activation of peripheral blood mononuclear cells from patient by epitope of infectious agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antigenic peptide, human papillomavirus; MAGE gene; BAGE-1 peptide; P18; human immunodeficiency virus; cancer antigen; tyrosinase; signal protein; anthrax lethal factor; LF; toxin; cationic fusion peptide; translocation; gene therapy; polycationic affinity handle; therapeutic protein; LFN.
                                                                                                                                            Human immunodeficiency virus type-1; HIV-1; T cell response; detection; peripheral blood mononuclear cell; PBMC.
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                                                                                                       HIV epitope env P18-IIIB amino acid residues 315-329 of gp160.
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AAW10348 standard; peptide; 15 AA.
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nes 8; Conservative
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Human immunodeficiency virus

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Query Match Best Loca Matches

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This is the antigenic human immunodeficiency virus peptide P18. This antigenic compound can be introduced into the cytoplasm of a cell by a new method where the cell is contacted with a fusion molecule comprising a delivery molecule. The delivery molecule can either be a polycationic affinity handle, LFN (the protective antigen binding domain of anthrax compound is linked to either of the delivery molecule related to LFN. The antigenic compound is linked to either of the delivery of the antigenic compound compound is linked to either of the delivery of the antigenic compound into a cell. The anthrax toxin system of the invention eliminates the need to generate fusion proteins with a toxin B molety, which alleviates problems associated with incorrect folding of lengthy fusion proteins. Small cationic fusion peptides substituted for LFN may reduce the possibility of steric interference with the biological activity of the translocated protein. The method is used for the introduction of antigens, e.g. MHC class I antigens or any other therapeutic protein.
                                                                                                                                                                                                                                                                    Introducing therapeutic proteins, especially antigens, into cells - using toxin molecules and/or polycationic handles for delivery.
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                                                                                                                                         Ballard JD,
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                                                                                                                                   Blanke SR, Milne JC, Lyszak EL,
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95US-0008518P.
96US-0019275P.
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                                                                                                                                                                                          Peptides AAW39275-W39300 are used in a novel method for inducing cytocoxic T-cell (TCC) activity specific for a broad array of HIV-1 separates using hybrid synthetic peptides. The method involves first administering a recombinant viral vector expressing the HIV-1 gp160 envelope glycoprotein and then administering at least 1 chimeric envelope glycoprotein and then administering at least 1 chimeric sequences corresponding to amino acid synthetic polypeptides having sequences corresponding to amino acid 315-329 of the gp160 envelope glycoprotein of HiV-1 strain IIIB, in which amino acid 325 is substituted by the corresponding amino acid from other strains or isolates, are used, a CTC response to a broad range of HIV-1 isolates can be elicited. These synthetic peptides are useful as vaccines for the prophylaxis or immunotherapy of HIV-1 virus infection
                                                                                                                        Inducing cytotoxic T-cell response to HIV - by administering gpl60 vector and chimeric gpl60 peptide(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                against human immunodeficiency virus - induces immune response
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gape
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                                                                    Germain RN, Berzofsky JA, Takahashi H;
                                        (USSH ) US DEPT HEALTH & HUMAN SERVICES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV-1 IIIB gp120 peptide fragment.
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                                                                                                                                                                    Example 1; Col 3; 25pp; English
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              91US-00760530
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 88US-00148692
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Matches 8; Conservative
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                                                                                               MPI; 1998-119931/11.
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26-JAN-1988;
               18-SEP-1991;
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23-JUN-1998
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This sequence represents a fragment of the human immunodeficiency virus (HIV) Type 1 strain IIIB gp120 protein. This sequence is used in a method

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resulting in the production of a vaccine against HIV which induces an immune response to the V3 epicope of HIV. This method which comprises the transplantation of an epitope of HIV at plural sites in the hypervariable region of immunoglobulin, the preparation of the epitope molecule histocompatibility antibody, and optionally chemically cross linking the epitope. An epitope histocompatibility antibody is also described in the specification which specifically responds to HIV, prepared by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
                                                                                                                                                          transplantation of an epitope comprising a peptide obtained from at least one V3 sequence of HIV. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (IIV) gp120 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gp120 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemocherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gp120, so prevents or inhibits 'bystander' apoptosis of uninfected T cells that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophilaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                      Gape
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100.0%; Pred. No. 0.1
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are bound to gpl20 protein, maximising induction of protective antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence is an epitope used in the construction of a novel fusion humanojobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, are bound to gpl20 protein, maximising induction of protective antibits bystander apported T cells that I cell immunity
                                                                                                                                                                                                                                                                                                                                                                                        B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                      Gaps
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                                                                                   Score 39; DB 2; Length 15;
Pred. No. 0.13;
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100.0%; Pred. No. v...
0; Mismatches
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                                                                               Query Match
Best Local Similarity
Matches 8; Conserv
                  cell immunity
                                                 Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                             25-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9836087-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-AUG-1998.
                                                                                                                                                                                                                                                                                             AAW76898;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scott D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sp.
                                                                                                                                                                                                                              RESULT 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus
                                                                                                                                                                                                                                                  AAW76898
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Score 39; DB 2; Length 15; Pred. No. 0.13;

100.0%;

Query Match Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence is an epitope used in the construction of a novel fusion human, ight chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tockrance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, are bound to gpl20 protein, maximising induction of protective antiviral T cell immunity
                                                                                                                                                                                                                                             B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                              Fusion immunoglobulin heavy chain HIV gp120 T cell epitope #19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 11; Page 120; 154pp; English.
                                                                                                              AAW76900 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (AMNA-) AMERICAN NAT RED CROSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                               98WO-US002766.
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                                                                                                                                                                               25-JAN-1999 (first entry)
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RAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zambidis E;
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Best Local Similarity
8; Conserv?
 RAFVTIGK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                               WO9836087-A1.
                                                                                                                                                                                                                                                                                                                                                              sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-FEB-1998;
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                                                                                                                                                AAW76900;
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                                                                               RESULT 52
                                                                                                                                                                                                                                                                                                                                                                  Ношо
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Sequence 15 AA;
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      29-SEP-1997;
12-DEC-1997;
                                                                                 or influenza.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JAN-1998;
                                          Letvin NL,
                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                      AAY24466;
                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kim T,
                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                       RESULT 55
                                                                                                                                                                                                                                                                                                               AAY24466
ID AAY2
                                                                                                                                                                                                                                                                                                                              ð
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                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                         Peptides AAW54903-W54941 can be used as an immunoadsorbent in an immunoassay for detecting antibodies to HIV gpl20, or as an immunogen for eliciting antibodies to HIV in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fusion protein; vaccine; cytokine; immunoglobulin; autoimmune disease; infectious disease; inflammatory disease; neoplastic disease; cancer; immune response; malaria; tuberculosis; hepatitis; AIDS; influenza; interleukin; IL-2; IG.
                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                         Immunoadsorbent; immunoassay; HIV gp120; immunogen; antibody; Human.
                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                     gp120 peptides - useful as immunoassay reagents or vaccine
                                                                                                                                                                                                                                                                                                                       100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.13;
                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                          gp120 envelope protein, peptide 127, analogue 127g'
                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                           Example 8; Column 21/22; 34pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence of gp120IIIB P18 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                         AAY06896 standard; peptide; 15 AA.
             AAWS4929 standard; peptide; 15 AA
                                                                                                                                                          91US-00663262.
91US-00726605.
94US-00326676.
                                                                                                                                                                                         (UNBI-) UNITED BIOMEDICAL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-US020321.
                                                                                        Human immunodeficiency virus
                                                                                                                                    95US-00488252
                                                                                                                                                    88US-00155321
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                                          (first entry)
                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                     1 RAFVTIGK 8
                                                                                                                                                                                                                       WPI; 1998-347301/30
                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                        Sequence 15 AA;
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                                                                                                                                    07-JUN-1995;
                                                                                                                                                           01-MAR-1991;
09-JUL-1991;
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                                           25-SEP-1998
                                                                                                                                                                           19-OCT-1994;
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                                                                                                                                                    12-FEB-1988;
                                                                                                       US5763160-A
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                                                                                                                     09-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                         AAY06896;
                            AAW54929;
                                                                                                                                                                                                        Wang CY;
                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                     HIV
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The invention relates to vaccine compositions comprising a vaccine and a timed-release formulation of a cytokine or cytokine/immunoglobulin fusion protein or plasmid The formulation or device releases the cytokine administration relamid at one or more temporal points subsequent to vaccine administration. The vaccines can be used for treating an autoimmune disease, an inflammatory disease, a neoplastic disease, or an immunologic disease in an individual. The vaccines can be used to elicit immune responses against diseases such as AIDS, malaria, tuberculosis, hepatitis C, hepatitis B, cancer or influenza. The methods can provide for enhancement of one or more immunologic parameters such as an antibody response, a cellular proliferative response as well as cytotoxic T-lymphocyte levels. In addition the Ig can increase the circulating half life of the cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                   New vaccine compositions for treating AIDS, maleria, tuberculosis, cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis B virus, HBV; X protein, cytotoxic T lymphocyte, liposome, CTL, antigen, immunity, liver cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Choi M,
                                                                                                   (BETH-) BETH ISRAEL DEACONESS MEDICAL CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MOGA-) MOGAM BIOTECHNOLOGY RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY24466 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; Page 22; 66pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
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97US-0060338P
97US-00990180
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The present invention describes peptide antigens AAY24459 to AAY24463 derived from the X protein of hepatitis B virus (HBV) which are recognized by cytcoxic T lymphocytes (CTL). The peptide antigens derived from HBV X protein are useful for inducing CTLs against the virus or inducing immunological tolerance to the virus, pH-sensitive liposomes containing the peptide antigens are used to induce cellular immunity so that CTLs specific to the virus can be produced. This is useful for prevention and treatment of HBV-associated diseases, especially HBV-associated liver cancer. pH-sensitive liposomes permit the selective transportation of anti-cancer drugs. The present sequence represents a
                                                                                                                                                                                                                                                                                                                                                    peptide used in an example from the present invention
                         Example 5; Page 14; 33pp; English
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 AA;
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100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.13; ive 0; Mismatches 0; Indels Ouery Match Ouer Local Similarity 100.00 Defines 8; Conservative 8 RAFVTIGK 15 1 RAFVTIGK 8 ò g

RESULT 56

AAY25189 standard; peptide; 15 AA. AAY25189; AAY25189 

HIV protein gp160 peptide fragment #1.

(first entry)

03-SEP-1999

Heat shock protein; HSP; complex; denatured protein matrix; antigen; vaccine; allergic disease; treatment; susceptibility; Th2; skin rash; allergic reaction; asthma; gp160.

Human immunodeficiency virus.

WO9929182-A1

17-JUN-1999.

98WO-US025734. 04-DEC-1998;

97US-00985548. 97US-00986234. 05-DEC-1997; 05-DEC-1997;

(UYNE-) UNIV NEW MEXICO STATE

Wallen ES, Moseley PL;

WPI; 1999-394912/33

Synthesizing heat shock protein complexes using a denatured protein

This invention describes a novel method for synthesizing heat shock protein (HSP) complexes comprises adding a heat shock protein to a denatured protein matrix for binding, and adding a complexing solution comprising a peptide to elute a heat shock protein-peptide complex. A HSP-antigen complex is useful as a vaccine for treating an allergic disease (in a mammal, preferably a human) to reduce susceptibility of the Th2 response, the complex comprising a HSP-antigenic peptide complex. The complex is administered to prevent a mammal from having an allergic reaction to an allergic disease, or administered to a mammal having an allergic disease, include asthma and skin rashes. Prior art methods or preventing/treating Example 1; Fig 1A; 33pp; English,

allergic diseases include antihistamines which treat only the symptoms, cortilosteroids which have severe side effects and desensitization therapy which has limited uses. The new method also allows more flexibility of use of peptide-based vaccines, as prior art HSP-based vaccines require isolation from a portion of the tumour itself. This sequence represents a peptide fragment from the HIV gp160 protein which is used in the method of the invention

Sequence 15 AA;

888888888

Gaps ö h Similarity 100.0%; Score 39; DB 2; Length 15; Similarity 100.0%; Pred. No. 0.13; 8; Conservative 0; Mismatches 0; Indels Query Match Best Local Similarity Best Loc Matches

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8 RAFVTIGK 15 œ 1 RAFVTIGK

a 8

RESULT 57

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Gaps

.; 0

AAY25204 standard; peptide; 15 AA

AAY25204;

03-SEP-1999 (first entry)

HIV V3 peptide fragment #10.

Heat shock protein; HSP; complex; denatured protein matrix; antigen; vaccine; allergic disease; treatment; susceptibility; Th2; skin rash; allergic reaction; asthma; V3 protein.

Human immunodeficiency virus.

WO9929182-A1

17-JUN-1999.

98WO-US025734 04-DEC-1998;

97US-00986234. 97US-00985548 05-DEC-1997; 05-DEC-1997;

(UYNE-) UNIV NEW MEXICO STATE.

Wallen ES, Moseley PL;

WPI; 1999-394912/33

Synthesizing heat shock protein complexes using a denatured protein matrix

Example 1; Fig 1B; 33pp; English.

protein (HSP) complaxes comprises adding a heat shock protein to a comprising and adding a complexing solution computed protein matrix for binding, and adding a complexing solution comprising a peptide to elute a heat shock protein-peptide complex. A HSP cartigen complex is useful as a vaccine for treating an allergic disease (in a mammal, preferably a human) to reduce susceptibility of the Th2 response, the complex comprising a HSP-antigenic peptide complex. The complex is administered to prevent a mammal from having an allergic disease, to reduce the allergic reactions. Allergic diseases conclude asthma and skin rashes, prior art methods or preventing/treating include asthma and skin rashes. Prior art methods or preventing/treating conclude asthma and skin rashes. Prior art methods or preventing/treating corticosteroids which have severe side effects and desensitization therapy which has limited uses. The new method also allows more therapy which has limited uses. The new method also allows more carticosteroids a limited uses. The new method also allows more carcine is sequence represents a peptide-based vaccines, as prior art HSP-based vaccines require isolation from a portion of the tumour itself. This caped in the method of the invention This invention describes a novel method for synthesizing heat shock 

(first entry) (revised)

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HIV-1 gp120 monoclonal antibody BAT123 residue 308 to 322.
                                                                  AAW72821 standard; peptide; 15 AA
                                                                                                                                                         17-OCT-2003
13-JAN-1999
                                                                                                            AAW72821;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents a HIV-1 cluster peptide vaccine conjugate (CLUVAC) sequence. The invention relates to a method for inducing a protective mucosal cytotoxic Tlymphocyte (CTL) response in a mammalian subject, which comprises contacting a mucosal tissue of the subject with induce a protective mucosal CTL response in a subject. The method can induce a protective mucosal CTL response in a subject. The method can used for protection against e.g. hepatitis A virus, papilloma virus, feline immunodeficiency virus, feline leukaemia virus, Listeria monocycogenes, M. tuberculosis, M. leprae, or Giardia lamblia. The method induces long-lasting protective mucosal immune responses. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1; CLUVAC; cluster peptide vaccine construct; cytotoxic T lymphocyte; protective mucosal CTL response; hepatitis A virus; papilloma virus; belane immunodeficiency virus; feline leukaemia virus; M. tuberculosis; Listeria monocytogenes; M. leprae; Giardia lamblia; immune response induction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Method for inducing a protective mucosal cytotoxic T lymphocyte immune
                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                  DB 2; Length 15;
                                                                                                            0; Indels
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                                                                                       0.13;
                                                                                                            0; Mismatches
                                                               100.0%; Score 39;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV-1 CLUVAC peptide, SEQ ID NO. 15.
                                                                                                                                                                                                                                                                                                                AAY05356 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; Page 85; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1.
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98US-0074894P
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                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                       Local Similarity 100.
1es 8; Conservative
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Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
                                                                                                                                                         1 RAFVTIGK 8
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                       Sequence 15 AA;
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17-FEB-1998;
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29-JUN-1999
                                                                                                                                                                                                                                                                                                                                                             AAY05356;
                                                                  Query Match
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RAFVTIGK 15

RAFVTIGK 8

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The present invention describes an antibody conjugate comprising an antibody (Ab) which binds to an epitope within amino acid residue 308-322 of gp120 and neutralises HIV-1, conjugated with a cytotoxic agent, an anti-viral agent or an agent which facilitates passage through the blood brain barrier. Also described is an antibody conjugate as above but where the Ab binds to an epitope within amino acid residue 298-312 of gp12 which neutralises HIV-1. The present sequence represents an HIV-1 gp120 monoclonal antibody BAT121 residue 308 to 322 from an example of the concern invention. The Ab are monoclonal Ab which bind to the gp120 protein on the envelope of HIV-1. They inhibit the infection of T-cells and also inhibit syncytium formation. The antibodies are group specific and also inhibit syncytium formation. The antibodies are group specific and also inhibit syncytium formation. The antibodies are group specific and also inhibit syncytium formation. The antibodies are group specific and also files of use a variety of uses, including the treatment and prevention of AIDS and AIDS related complex. They are especially used to kill infected T-cells. (Updated on 17-OCT-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibody conjugate comprising monoclonal antibody - which binds to epitope within amino acid residue of gpl20 which neutralises HIV-1 conjugated with, e.g. cytotoxic agent.
HIV-1, gpl20, epitope, monoclonal antibody, envelope, neutralise, inhibit, infection, T-cell, inhibit syncytium formation, AIDS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chang T;
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                                                                                                    Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                    87US-00057445.
87US-00137861.
89US-00343540.
92US-00895197.
                                                                                                                                                                                                                                                                                            93US-00026276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TANO-) TANOX BIOSYSTEMS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SC, Kim YW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-008810/01.
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Best Local Similarity
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25-APR-1989;
                                                                                                                                                                US5834599-A.
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98WO-US015580
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Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                               Rakowicz-Szulczynska EM,
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                                                                                                                                                                               29-JUL-1998;
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05-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents an epitope of the gp120 protein of human immune deficiency virus (HIV)-1. The sequence comprises amino acids 308 to 322 of gp120. The specification describes monoclonal antibodies which bind to epitopes of the gp120 protein. Specifically, these antibodies are designated BAT123, 267 and 085. Monoclonal antibodies neutralise HIV-1, used to treat acquired immune deficiency syndrome (AIDS) and AIDS-related complex, by passive immune deficiency syndrome (AIDS) and AIDS-related agents, and in extracorporeal systems. They can also be used as agents, and in extracorporeal systems. They can also be used as screen for neutralising epitopes, potentially useful in vaccine development. (Updated on 20-MAR-2001 to correct PR field.) (Updated on 17-0CT-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal antibodies directed against regions of gp120 of human immune deficiency virus-1 - are neutralising and able to inhibit infection of cells and formation of syncytia, used for treatment, prevention or diagnosis of acquired immune deficiency syndrome.
                                                                                                            Epitope, gpl20 protein, monoclonal antibody, HIV-1, antibody BAT123, antibody BAT267, antibody BAT085, T cell infection inhibition, syncytia formation, acquired immune deficiency syndrome, AIDS, AIDS-related complex, passive immunisation, antiviral, cytotoxic,
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                                                                        Epitope of HIV-1 gp120 protein which binds antibody BAT123.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chang TW;
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                                                                                                                                                                                       viral load measurement; vaccine
                                                                                                                                                                                                                               Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sun CRY,
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91US-00767533.
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                                      (first entry)
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                            22-SEP-1992;
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                                                                                                                                                                                                                                                                                                                                                                                    29-MAY-1987;
17-OCT-2003
                  20-MAR-2003
03-MAR-1999
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22-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fung MSC,
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HIV-1 gp120 amino acids 308-322.

RESULT 61
AAY04680
ID AAY0
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AC AAY0
XX
DT 17-C
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XX

Matches

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This peptide corresponds to amino acids 308-322 from the gp120 protein of the human immunodeficiency virus type 1 (HIV-1). The peptide is used to generate the monoclonal antibody Mab 5023 The invention relates to the use of homology between HIV-1 antigens and breast, gynecological and prostate cancer antigens to develop agents for use in the detection and treatment of such cancers. The method especially uses an antibody which recognises the p160, p80, p45 and p24 cell membrane proteins and the p24 chromatin protein. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of HIV-1 polypeptides - for developing products for the detection and treatment of breast, gynecological and prostate cancers.
gpl20; HIV-1; monoclonal antibody; homology; antigen; breast; prostate;
gynecological; cancer; detection; diagnosis; cell membrane; chromatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunogen, particulate composition; immune response; assessment; target skin site; skin immune reaction; HIV-1; immunocompetence; antibody; cell mediated immunity; antigen exposure; allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.13; Live 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                   (RAKO/) RAKOWICZ-SZULCZYNSKA E M.
                                                                   Human immunodeficiency virus 1.
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99US-0139045P.
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10-JUN-1999;
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The invention relates to a method of using an immunogenic compound from a selected agent in the manufacture of a particulate composition for assessable an immune response against the selected agent in an individual. The method comprises: (a) accelerating the particulate composition into a target skin site in the individual; and (b) assessing the target site to determine the presence or absence of a localized skin immune reaction, where the presence of the immune reaction is indicative of an immune response against the selected agent. Peptides AAV83316-Y83325 represent buman immunodeficiency virus type 1 (HIV-1). The method is used to detect assessing immunocompetence, antibody and cell mediated immunity, antigen examples.
                                                                        comprises accelerating a particulate composition, containing an immunogenic compound from a selected agent, into the target skin site of the individual.
                                                                 Assessing an immune response against a selected agent in an individual
                                                                                                                                                                              Disclosure; Page 23; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2003 to standardise OS field)
                    WPI; 2000-257072/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 AA;
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100.0%; Score 39; DB 3; Length 15; 100.0%; Pred. No. 0.13; 0; Mismatches 8; Conservative RAFVTIGK 15 RAFVTIGK 8 셤 ઠે

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Gaps

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0; Indels

AAY66439 standard; peptide; 15 AA. AAY66439; RESULT 63 **AA**X66439 

(first entry) (revised) 12-SEP-2003 22-FEB-2000 HLA-A2-binding HIV-1 GP120 CTL epitope #241.

HIV-1; MHC; major histocompatibility complex, Class I; HLA-A2; human leukocyte antigen; CTL; cytotoxic T-cell; epitope; allele; binding; conserved; genome; peptide; targetting; toxic; drug; antibody; antigen; antiviral; molecular conjugate therapeutic; diagnosis; treatment; pathogen; localisation; quantification; detection; infection; drug resistance; immune response

Human immunodeficiency virus 1.

WO9949893-A1

07-OCT-1999.

99WO-US007111 31-MAR-1999; 98US-00052530. 31-MAR-1998;

(UYBO-) UNIV BOSTON

Zhang Weng Z, Vaccaro D, Gulukota K, Berzofsky J, Delisi C,

ΰ

WPI; 2000-038361/03

Novel methods for designing molecular conjugate therapeutics which are used for diagnosis, imaging and treatment against pathogens.

Example 3; Page 50; 62pp; English

99WO-US007111. 31-MAR-1999; 98US-00052530. 31-MAR-1998;

(UYBO-) UNIV BOSTON

Zhang C; Weng Z, Gulukota K, Vaccaro D, Delisi C, Berzofsky J,

WPI; 2000-038361/03.

AAY66421-Y66453 are cytotoxic T-cell epitopes derived from conserved portions of the HIV-1 genome that are presented by HIA-A2 MHC (major histocompatibility complex) Class I molecules. The peptides are used to construct targetting antigens complising one or more peptides bound to the corresponding MHC Class I molecule, which can be used to raise antibodies. The antibody may then be used as a targetting vehicle to the corresponding MHC Class I molecule, which may result in adverse side affects. The invention relates to improved methods for the design of molecular conjugate therapeutics for the diagnosis and treatment of infections caused by pathogens with a high mutation rate (such as HIV-1). The method involves identifying conserved peptide-encoding regions among the genomes of multiple variants of a pathogen, identifying the Class I MHC molecules which occur with greatest frequency in a population of the corresponding Class I MHC molecules. The MHC-binding peptides and to the Class I MHC molecules are selected and used to construct targetting antigens, which are in turn used to produce the construct targetting antigens, which are in turn used to produce construct targetting antigens, which are in turn used to produce targetting antigens, which are in turn used to produce targetting antigens, which are in turn used to produce targetting antigens, which are in turn used to produce targetting antigens, which are in turn used to produce targetting and so to detect and treat viral infection. The methods of the invention mitigate against the development of viral resistance to drugs and to the immune response, as well as providing a solution for acressible to x colls. In addition, the methods eliminate the virus, whereas current therapies only arrest viral infection. (Updated on 12-CNE SEP-2001 to standardise OS field) ö HIV-1; MHC; major histocompatibility complex; Class I; Caucasoid; HLA; human leukocyte antigen; CTL; cytotoxic T-cell; Caucasian; epitope; allele; binding; conserved; genome; peptide; targetting; toxic; drug; antibody; antigen; antiviral; molecular conjugate therapeutic; diagnosis; treatment; pathogen; localisation; quantification; detection; infection; drug resistance; immune response. Gaps .; 0 100.0%; Score 39; DB 3; Length 15; 100.0%; Pred. No. 0.13; 0; Indels HLA-A3-binding HIV-1 GP120 CTL epitope #257. 0; Mismatches AAY66455 standard; peptide; 15 AA. Human immunodeficiency virus 1. (first entry) 8; Conservative (revised) 8 RAFVTIGK 15 œ Local Similarity 1 RAFVTIGK Sequence 15 AA; WO9949893-A1 12-SEP-2003 22-FEB-2000 07-OCT-1999 AAY66455; Query Match Matches RESULT 64 AAY66455 ઠે 셤

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ANY 66454-Y66458 are cytotoxic T-cell epitopes derived from conserved portions of the HIV-1 genome that are presented by MHC (major continues of the HIV-1 genome that are presented by MHC (major compatibility complex) Class I alleles found with high frequency among Caucasoids in the USA. The peptides are used to construct targetting antigens comprising one or more peptides bound to the corresponding MHC class I molecule, which can be used to raise antibodies. The antibody may then be used as a targetting vehicle to deliver a potentially toxic drug to its target site of action, rather can deliver a potentially toxic drug to its target site of action, rather confugate therapeutics for the diagnosis and treatment of infections caused by pathogens with a high mutation rate (such as HIV-1). This method involves identifying conserved peptide-encoding regions among the genomes of multiple variants of a pathogen, identifying the Class I will method involves identifying conserved peptide-encoding regions among the genomes of multiple variants of a pathogen, identifying the Class I conserved peptides bind to the Class I MHC molecules. The MHC-binding peptides and interest terresponding Class I MHC molecules are selected and used to construct targetting antigens i MHC molecules are selected and used to construct targetting antigens, which are in turn used to produce targetting antibodies. The methods may be used in localisation, quantification and in situ detection of specific peptide-MHC class I complexes and also to detect and treat viral infection. The methods of the invention mitigate against the development of viral resistance to drugs and to the immune response, as well as providing a solution for targetting taxic compounds to destery virals sequestered in sites not accessible to T cells. In addition, the methods sequestered in 12-construct therapies only arrest viral replication. (Updated on 12-construct therapies only arrest viral replications.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunogenic particle; human immunodeficiency virus; HIV; cytostatic; antiarthritic; antiinflammatory; cell-mediated immune response; cancer; rheumatoid arthritis; inflammatory disorder; viral infection.
                     Novel methods for designing molecular conjugate therapeutics which are used for diagnosis, imaging and treatment against pathogens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 39; DB 3; Length 15; 100.0%; Pred. No. 0.13; 0; Indels ive 0; Mismatches 0; Indels
                                                                                         Example 3; Page 51; 62pp; English.
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Best Local Similarity 100.0%;
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(first entry)
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01-FEB-2001
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This invention relates to artificial immunogenic particles comprising glycolipids having a lipidic and a saccharide portion and peptide-lipid conjugates having a lipidic and a peptide portion embedded into a lipid based matrix. The peptide portion of the particle may be of viral origin. Peptides AAY85579-Y85591 are human immunodeficiency virus (HIV) related particles which can be used as the peptide portion in an immunogenic particles which can be used as the peptide portion in an immunogenic and antiinflammatory activity. The immunogenic particles are used for inducing a cell-mediated immune response in a host directed towards the peptide portion of the peptide-lipid conjugate. This means that the particles may be used to treat diseases such as cancer, rheumatoid arthritis, inflammatory disorders or viral infections such as HIV. (Updated on 12-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New peptide 3, amide and heterocyclic compounds and saccharide conjugates used for inhibiting chemokine induced activity and for treating e.g. stroke, vascular diseases, autoimmune diseases and tumor growth.
                                                                                      Lipid based artificial particles useful for inducing a cell mediated immune response in a subject having cancer, comprises a lipid based matrix, glycolipids and peptide-lipid conjugates embedded in the matrix.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Macrophage recruitment; chemokine derivative; MCP-1; osteoporosis; monocyte chemoattractant protein-1; inflammation; atherosclerosis; HIV; AIDS; stroke; psoriasis; autoimmune disease; hypertension; endoroxaemia; basophil-mediated disease; myocardial infarction; acute ischaemia; rheumatoid arthritis; contraception.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 39; DB 3; Length 15; 100.0%; Pred. No. 0.13; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human chemokine derived peptide #27.
               Sugimoto M, Arella M, Furuichi Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB15875 standard; peptide; 15 AA.
                                                                                                                                                                 Claim 10; Page 34; 39pp; English.
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99US-00271192.
99US-00452406.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.
Best Local Similarity 100.
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8 RAFVTIGK 15
                                                     WPI; 2000-664891/64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200042071-A2
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01-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 66
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(SAPI-) SAPIENTIA THERAPEUTICS LTD. (AGEN-) AGENE RES INST CO LTD.

23-MAR-2000; 2000WO-CA000319

99US-00276057

25-MAR-1999;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a pharmaceutical compound (I) that contains an N-terminal glutamic acid (Glu) or glutamine (Gln) residue in the form of an addition salt with a strong, physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition containing at least one (I); (b) a vaccine containing at least one (I)
                                                                                                                                                                                                                                autoimmune diseases, atheroscierosis, osteoporosis, HIV infection and AIDS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotoxaemia, myocardial infarction, acute ischaemia and rheumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in gene therapy for the same diseases, as well as in the production of
                                                                                                              chemokines which are used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences AAB1595-B15794, AAB15803-B15813 and AAB15831-B15848. These chemokine derivatives can be used to treat diseases such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Сарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stabilized pharmaceutical containing N-terminal glutamic acid or glutamine, useful e.g. in anti-melanoma vaccines, is an addition salt with strong acid.
                                                                                        present invention concerns the identification of a number of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 39; DB 3; Length 15; 100.0%; Pred. No. 0.13; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vaccine related MHC ligand peptide SEQ ID NO:186.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Klinguer-Hamour C, Corvaia N, Beck A,
                            Disclosure, Fig 18; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (FABR ) FABRE MEDICAMENT SA PIERRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAM99083 standard; peptide; 15 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-MAR-2001; 2001WO-FR000872.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200170772-A2
                                                                                                                                                                                                                                                                                                                                                                                                                          animal models
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07-DEC-2001
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where this is a major histocompatibility complex (WHC) ligand (Ia); (c) a method for in vitro diagnosis of diseases associated with the presence of (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process for preparing (I). (I) has immunomodulator, endocrine, antiallergic, neuroprotectant, virucidal, bactericidal, antiparasitic, fungicidal and corporate activities. (I) are useful, in human or veterinary medicine, in pharmaceutical compositions (for treating immune disorders, e.g. in munne deficiency, autoimmune states, hypersensitivity, allergy, graft cefection, infection, hormonal disorders and central nervous system cliented in there (I) is a MMC ligand (Ia), in vaccines for treatment or prevention of: (i) viral, bacterial, parasitic or fungal infections; or (ii) of cancers A particular application is in anti-complexed with interactions between MMC and (I), e.g. melanoma and human immunoma infection. AMMSSSS to present peptides which can be used in pharmaceutical compounds from the present invention. (Updated on 11-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protection, endogenous therapeutic peptide, peptidase, conjugation, blood component, modification, succinimidyl, maleimido group, amino, hydroxyl; thiol; hormone; growth factor, neurotransmitter.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 39; DB 4; Length 15; 100.0%; Pred. No. 0.13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Virus related peptide SEQ ID NO:1521.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB92345 standard; peptide; 15 AA.
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99US-0153406P.
99US-0159783P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7 RAFVTIGK 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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AAB92345
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(I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half intracellular upsake and interference with physiological processes. AAB90829 to AAB9241 represent invention Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter. Thibaudeau K; 100.0%; Score 39; DB 4; Length 15; 100.0%; Pred. No. 0.13; tive 0; Mismatches 0; Indels Virus related peptide SEQ ID NO:1524. AAB92348 standard; peptide; 15 AA 99US-0153406P. 99US-0159783P. 99US-0134406P. 17-MAY-2000; 2000WO-US013576 (first entry) 8; Conservative (CONJ-) CONJUCHEM INC. 1 RAFVTIGK 8 Local Similarity Sequence 15 AA; WO200069900-A2 Homo sapiens 17-MAY-1999; 22-JUN-2001 10-SEP-1999; 15-OCT-1999; Synthetic. ထ AAB92348; Query Match Best Loc Matches RESULT 69 AAB92348 88666666666668888 ò 음

Bridon DP, Ezrin AM, Milner PG, Holmes DL, WPI; 2001-112059/12. Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity. Disclosure; Page 703; 733pp; English.

The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attracted to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or

reduces the action of peptidases to increase length of activity (half life) and specificity as bonding to large molecules decreases intracellular uptake and interference with physiological processes. AAB90829 to AAB92441 represent peptides which can be used in the exemplification of the present invention 88888888

Sequence 15 AA;

Gaps ö Score 39; DB 4; Length 15; Pred. No. 0.13; 0; Mismatches 0; Indels 100.0%; 8; Conservative Query Match Best Local Similarity Matches 8; Conserv

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AAB68601 standard; peptide; 15 AA. (revised) 11-SEP-2003 25-APR-2001 AAB68601; RESULT 70 AAB68601

HIV gp120 V3 loop peptide #1. (first entry)

HIV gp120 V3 loop; liposome composition; HIV infection. Human immunodeficiency virus 1

US6180134-B1.

95US-00480332. 07-JUN-1995; 23-MAR-1993; 30-JAN-2001.

93US-00035443. (SEQU-) SEQUUS PHARM INC. 29-SEP-1994; 

Woodle MC, Martin FJ, Barenholz Y; Zalipsky S,

WPI; 2001-201897/20.

Liposome composition for use in treating septic shock comprises liposomes having an outer surface layer of polyethylene glycol chains, and a polypeptide or polysaccharide effector molecule.

Disclosure, Fig 13; 32pp; English.

in present inventors to a injustment composition computation composition tender surface layer of polyearchylene glycol chains, each having a free distal end. A polypeptide or polysaccharide effector cach having a free distal end. A polypeptide or polysaccharide effector molecule is covalently attached to a portion of the distal ends. The effector interferes with specific binding of pathogen or cell in a bloodstream when administered in free form. The liposome composition may be used in treating a condition mediated by binding a pathogen or cell in the bloodstream, to a target cell or cell carrix. It can be used in treating septic shock, toxic shock, colonic inflammation, leukaemic cell proliferation, or HIV infection. The present sequence is a peptide of the V3 loop of HIV envelope protein gpl20 binds to the CD4 receptor during HIV infection of lymphocytes. By introducing the present peptide, the CD4 receptors are blocked, thereby introducing HIV infection of lymphocytes. By introducing HIV infection of lymphocytes. By introducing HIV infection of lymphocytes by introducing the present peptide, the CD4 receptors are blocked, thereby introducing HIV infection of lymphocytes are blocked, thereby introducing HIV infection of lymphocytes are blocked, thereby introducing the present peptide, the CD4 receptors are blocked, thereby introducing the present peptide on 11-SBP-2003 to standardise os The present invention relates to a liposome composition comprising

Sequence 15 AA;

8 RAFVTIGK 15

RESULT 72

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The invention relates to a composition comprising an expression vector bound to an aggregated protein-polycationic polymer conjugate or auspension. The expression vector contains a promoter polynucleotide sequence operatively linke to a polynucleotide sequence encoding an antigen which is a fragment of a gene or genome associated with an arthritis, vasculitis, and multiple sclerosis, pathogenic genomes consisting of bacterium, fungus, proteosa and virus such as human chimumodeficiency virus (HCV), herpes simplex virus (HCV), influenza and respiratory syncytial virus (RSV), hepatitis C virus (HCV), influenza and respiratory syncytial virus (RSV), or optionally comprising a nucleotide sequence encoding a cytokine (or a cytokine expression vector), is useful for inducing an immune response (systemic and/or mucosal) in an organism. The cytokine expression vector contains a sequence for granulocyte macrophage-colony stimulating factor (GM-CSF) or interleukin-12 (IL-12). The polynucleotide sequences encoding the antigen and the cytokine are under transcriptional control of same or with the companian contains as a contain as a cytokine are under transcriptional control of same or contains a contain as 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV; human immunodeficiency virus; cytostatic; immunosuppressive; pl8; virucide; antibacterial; fungicide; protozoacide; antirheumatic; vaccine; antiinflammatory; antiarthritic; neuroprotective; rheumatoid arthritis; cancer; multiple sclerosis; immune response; vasotropic; gene therapy; autoimmune disease; vasculitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Composition for oral delivery of vaccines, comprises expression vector containing antigenic genomic sequence, bound to aggregated protein-polycationic polymer conjugate or suspension.
                                                      Gaps
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100.0%; Score 39; DB 4; Length 15; 100.0%; Pred. No. 0.13;
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                                                         Mismatches
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                                                                                                                                                                                                                                                                                                   AAE15743 standard; peptide; 15 AA
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                                                   Conservative
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                                                                                                                                                               RAFVTIGK 15
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                                                                                                         1 RAFVTIGK 8
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Matches 8; Conser
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  Query Match
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AAE15743
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administering one or more virus like particles carrying an immunogen. It is also useful for administering to a patient for generating antibodies specific for one or more wirus like patient for generating antibodies and end or more immunogens, such as Blueongue virus (BTV), foot and mouth disease virus (FMDV), influenza virus and human immunodeficiency virus (HIV). The invention provides an effective means of delivering multiple peptide components representing viral/cumour antigenic groups to elicit protective immunity, which has not previously been possible. The present amino acid sequence represents one of a collection (AAUS6022-AAUS6045) of HIV epitopes that were used in the methods of the invention as immunogens. These epitopes were used to construct chimeric N31-TUBs (virus-derived tubules) which show immunogenicity. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a new vaccine comprising a non-
replicating vaccine delivery vehicle (which comprises a non-infectious
recombinant viral tubule) carrying one or more immunogens. The invention
is useful for inducing an immune response, preferably anti-viral, in a
subject. The administration of the vaccine is preferably followed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A vaccine, for inducing an antiviral immune response, comprises a non-replicating vaccine delivery vehicle (which comprises a non-infectious recombinant viral tubule) carrying one or more immunogens.
                                                                                                                                                                                                       Vaccine; non-replicating; viral tubule; immunogen; antibody; BTV; Bluecongue virus; foot and mouth disease virus; FMDV; influenza virus; human immunodeficiency virus; HIV; protective immunity; epitope; TUB; virus-derived tubule; anti-HIV; virucide.
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                                                                                                                                                               HIV epitope, HIV-1 gp120 H-2Dd,help, peptide sequence.
                 AAU96031 standard; protein; 15 AA.
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Query Match
100.0%; Score 39; DB 5; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.13;
Matches 8; Conservative 0; Mismatches 0; Indels

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  An adjuvant of a DNA vaccine composed of alpha-antigen genes derived from acid-fast bacterium.
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                                                                                                                                                                                                        Adjuvant; acid-fast bacterium; acquired immunodeficiency syndrome; MNA vaccinie, ALDS; hepatitis C virus; alpha-antigen gene; CTL; Mycobacterium kansasaii; antigenic; immunogenic; epitope; human immunodeficiency virus; HIV.
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                                     AAU97690 standard; peptide; 15 AA
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                                                                                                                                                                    HIV CTL epitope peptide sequence.
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                                                                                                             (revised)
(first entry)
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13-AUG-2002
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07-OCT-2002
                                                                          AAU97690;
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RESULT 73
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                   AAU97690
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Administering a non-viral vector encoding a co-stimulatory molecule alongside a peptide or protein T cell epitope, elicits increased response to the antigen and is useful to enhance peptide and protein based vaccines and treatments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method for eliciting an immune response in a subject. The method comprises administering a peptide or protein antigen comprising T-cell epitope(s) (e.g. tumour antigen, viral or non-viral antigen) coordinately with a non-viral vector comprising a polymucleotide encoding a T-cell co-stimulatory molecule. Viral peptide antigens may include human immunodeficiency Virus (HBV) antigen, virus (HBV). The method is useful to elicit an immune response in a subject, and to supplement and enhance peptide and protein based vaccines and treatment methods. ABGGGGGO-BORDON Cepresent HIV-1 peptide antigens useful in the method of the present invention. (Updated on 29-AUG-2003 to
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Eliciting an immune response, peptide antigen, T-cell epitope, tumour antigen, viral antigen, non-viral vector, HIV-1, T-cell co-stimulatory molecule, human immunodeficiency virus,
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                                                                                                                          Human immunodeficiency virus 1; (IIIB isolate).
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(BERZ/) BERZOFSKY J.
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                                                                           immunostimulant
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                                                                                                                                                                                                               The present invention relates to a method for eliciting an immune response in a subject. The method comprises administering a peptide or protein antigen comprising T-cell epitope [6] (e.g. tumour antigen, viral or non-viral antigen) coordinately with a non-viral vector comprising a polynucleotide encoding a T-cell co-stimulatory molecule. Viral peptide antigens may include human immunodeficiency virus (HIV) antigen, hepatitis B virus (HBV), herpes simplex virus (HSV) or human papilloma virus (HPV). The method is useful to elicit an immune response in a subject, and to supplement and enhance peptide and protein based vaccines and treatment methods. ABG68640-ABG68700 represent HIV-1 peptide antigens useful in the method of the present invention. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                           Administering a non-viral vector encoding a co-stimulatory molecule alongside a peptide or protein T cell epitope, elicits increased response to the antigen and is useful to enhance peptide and protein based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytolytic T lymphocyte; epitope; vaccine; prophylaxis; HIV infection; human immunodeficiency virus; acquired immune deficiency syndrome; CTL; gene therapy; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 39; DB 5; Length 15; 100.0%; Pred. No. 0.13;
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                                                                                                                                                                                             Disclosure; Page 7; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAE35161 standard; peptide; 15 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                POWDERJECT VACCINES INC. POWDERJECT RES LTD.
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18-MAY-2001; 2001US-0291655P.
                             14-MAR-2001; 2001US-00810310.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-MAY-2002; 2002WO-GB002336
                                                 15-MAR-2000; 2000US-0189396P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 100 les 8; Conservative
                                                                                                                                                                          vaccines and treatments.
                                                                                                    Khleif S, Berzofsky J;
                                                                   (KHLE/) KHLEIF S.
(BERZ/) BERZOFSKY J.
                                                                                                                                                                                                                                                                                                                                                                                                                                     RAFVTIGK 15
                                                                                                                                                                                                                                                                                                                                                                                                                  RAFVTICK 8
                                                                                                                       WPI; 2002-507231/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV CTL epitope #6.
                                                                                                                                                                                                                                                                                                                                                       Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200294313-A2.
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          18-APR-2002.
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AAE35161
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HIV; antigen; epitope; T cell; MHC; major histocompatibility complex; CTL; cytotoxic T lymphocyte; HIV infection; cancer; tuberculosis; tumour; hepatitis; melanoma; breast cancer; Hodgkin lymphoma; nasopharyngeal carcinoma; vaccine; immune response; hyaluronic acid; HA; CD8+ T cell; CD4+ T cell; viral infection; bacterial infection; tungal infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modulating an immune system response to an antigen in a mammal, comprises administering a particle-free therapeutic comprising a hyaluronic acid polymer analogue covalently linked to a peptide that comprises a T cell
                                                                                                                                                                                                                                                                                                                          The invention relates to a recombinant nucleic acid comprising a nucleic acid sequence encoding an antigen containing two or more cytolytic T lymphocyte (CTL) epitopes or its analogues. Sequences of the invention are used in vaccines and are useful for the treatment and prophylaxis of human immunodeficiency virus (HIV) infection, particularly acquired therapy. The present sequence is HIV CTL epitope. This sequence is used in the exemplification of the invention
                                                                                                              Recombinant nucleic acid for the treatment and prophylaxis of acquired immunodeficiency syndrome, comprises a nucleic acid sequence encoding an antigen containing two or more cytolytic T lymphocyte (CTL) epitopes or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gape
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 100.0%; Score 39; DB 6, Length 15; Similarity 100.0%; Pred. No. 0.13; 8; Conservative 0; Mismatches 0: Indela
Haynes J, Shipley T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADN14074 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                     Claim 1; Col 78; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus 1.
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                                                        WPI; 2003-148439/14.
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Best Local Similarity
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                                                                                                                                                                                                               its analogs
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Matches
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The invention relates to modulating an immune system response to an antigen in a mammal comprising administering to the mammal a particlefree therapeutic comprising a hyaluronic acid (HA) polymer analogue covalently linked to at least one peptide that comprises a T cell epitope comprises a sequence of at least about eight amino acids of the antigen. Also included are a method of improving major histocompatibility complex (MHC) presentation of a T cell epitope of an antigen in a mammal (comprising administering to the mammal the conjugate). The T cell epitope is recognised by a major histocompatibility complex (MHC) class I molecule and by a CD8+ T cell of the mammal. Or the mammal to an MHC class II molecule and a CD4+ T cell of the mammal.

The immune system response comprises a cytotoxic T lymphocyte, a CD4+T cell, or an antibody that recognises the antigen. The immune system response comprises the antigen of a pathogenic agent of a cusping sization of the conjugate, where the antigen is an antigen of a cumour cell. The immune system response to the antigen is an antigen of a cusping ration of the conjugate, where the antigen of a pathogenic agent or organ transplanted to the mammal. The composition and methods are useful for modulating, i.e. enhancing or diminishing, an immune system response to an antigen in a mammal. The composition of a crisque or organ transplanted to the mammal. The composition of a conjugates are useful as peptide vaccines against an antigen of conjugates are useful as peptide vaccines against an antigen of a conjugates are useful as peptide vaccines against an antigen, are useful for recenting or preventing diseases associated with any of the antigens above e.g. HIV infection, cancer, tuberculosis, hepatitis, antinomal parasticency. igens above e.g. HIV infection, cancer, tuberculosis, hepatitis, anoma, breast cancer, Hodgkin's lymphoma and nasopharyngeal carcinoma. peptide vaccine compositions of the present invention do not require than peptide vaccines of the prior art. The present sequence is an HIV-1 derived epitope suitable for the vaccine of the invention. additional adjuvants, but still induce a stronger cell-mediated response Gaps vaccine; virucide; antibacterial; immunosuppressive; antiallergic; ö 100.0%; Score 39; DB 7; Length 15; 100.0%; Pred. No. 0.13; ive 0; Mismatches 0; Indels Immune response induction composition peptide adjuvant #2. (UYPO-) UNIV POHANG SCI & TECHNOLOGY. (GENE-) GENEXINE CO LTD. ADR04041 standard; peptide; 15 AA 30-JAN-2004; 2004WO-KR000177 30-JAN-2003; 2003KR-00006393 cytostatic, peptide adjuvant (first entry) Local Similarity 100. 8 RAFVTIGK 15 1 RAFVTIGK 8 WO2004067020-A1 Sequence 15 AA; 04-NOV-2004 12-AUG-2004. Synthetic. Query Match ADR04041; Best Loc Matches ADROUAL 78
ADROUAL 1D ADROUAL
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Sung Y;

Kim Y,

Ryu S,

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Lee C,

Park S, Yang S,

Park K,

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                                                    encoding an immunogenic protein, useful for inducing immune responses against diseases e.g. HIV infection, autoimmune diseases, tuberculosis or
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                                                                                                                                                                       peptide adjuvant and a DNA vaccine encoding an immunogenic protein. The composition may also comprise a gene of the influenza virus, preferably the neuraminidase gene. The vaccine composition is useful for inducing immune responses against diseases comprising HIV infection, herpes simplex virus (HSV) infection, influenza virus infection, hepatitis A or a bifection, papallomavirus infection, tuberculosis, tumour growth, autoimmune diseases or allergies. The present sequence is a peptide adjuvant useful in the composition of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lipopeptide; lipoprotein; vaccine; cytotoxic T-cell; lymphocyte; HIV;
human immunodeficiency virus; AIDS; cancer; tumour cells; CB1; CB2; CB3.
                                    New vaccine composition comprising a peptide adjuvant and a DNA vaccine
                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                           The present invention relates to a vaccine composition comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lipopeptide(s) which stimulate cytotoxic T-cells - for treating HIV,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (INSP ) INST PASTEUR LILLE.
(INRM ) INSERM INST NAT SANTE & RECH MED.
(INSP ) INST PASTEUR.
                                                                                                                          Example 2; Page 21; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR24939 standard; protein; 16 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91EP-00403446.
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                                                                                                                                                                                                                                                                                                                                                                                                          8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV peptide ENV 312-327.
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WPI; 2004-580853/56
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                      Sequence 15 AA;
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09-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detection, purification and elimination of antigen-specific lymphocytes for producing cytotoxic T cells for immuno-therapy of cancers and viral infection.
peptide. The lipoproteins are useful in vaccines and acts by inducing cytocoxic T lymphocytes against the HIV virus antigen from which the peptide is derived. See also AAR24938 and AAR24940. (Updated on 25-MAR-2003 to correct PM field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antigen; major histocompatibility complex; MHC; lymphocyte; detection; immobilisation; cytotoxic T-cell; tumour; leukaemia; lymphoma;
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                                                                                                                                                       100.0%; Score 39; DB 2; Length 16; 100.0%; Pred. No. 0.14; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                        AAW68326 standard; peptide; 16 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MHC binding peptide Env.312-327.
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                                                                                                                                                                        Best Local Similarity
Matches 8; Conserv
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                                                                                                                Sequence 16 AA;
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14-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                            RESULT 80
AAW68326
ID AAW68326
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AAW68326
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AC AAW68.
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Score 39; DB 2; Length 16; Pred. No. 0.14;

100.0%;

Query Match Best Local Similarity

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The present invention describes a recombinant DNA molecule (I) containing a sequence (Ia) that encodes an altered MHC (major histocompatibility complex) Class I determinant (II) comprises a polypeptide with alphal, alpha2, alpha3 and betez-microglobulin domains, in which alpha1 and betez-microglobulin domains, in which alpha1 and beta2 care covalently linked, thorough C- and N-termini respectively, via a nucleotide spacer sequence encoding a polypeptide. (II) includes an carigon-binding site and when (II) and the antigen are associated they are recognized by a mammalian T cell receptor (TCR). (I) are used to produce (II) which are used to study functional interactions between the various MHC domains. They can also be used to modulate (in vivo or in vitro) the immune system by inducing an effector response (cytotoxicity, antibody synthesis, phagorytosis) of immune system cells, typically for treating, or immunising against; cancer, acquired immune deficiency syndroms, lupus erythematosus, multiple sclerosis, typically for treating, or immunising against; cancer, acquired immune deficiency capanosis, therapy, studying MHC-associated cellular chies, but also for selective destruction of autoreactive cells, cueful for diagnosis, therapy, studying MHC-associated cellular cyccesses and for affinity purification) AAZ57558 and AAY68186 to hyperical in the present cyconetion. (Updated on 12-SEP-2003 to standardise OS field)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New recombinant DNA encoding covalently linked form of major histocompatibility complex Class I determinant, used for immune system stimulation, e.g. for treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MHC class I; major histocompatibility complex; microglobulin; antigen; immune response; immunisation; AIDS; multiple sclerosis; toxic shock; cancer; lupus erythematosus; snake bite; cytostatic; antiviral; immunomodulatory; dermatological; immunosuppressive; antinflammatory;
   Gaps
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   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Altered MHC determinant binding peptide SEQ ID NO:35.
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   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Abastado J;
                                                                                                                                                                                                                                                                                                                   AAY68203 standard; peptide; 16 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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   8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kourilsky P, Mottez E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PASTEUR
                                                                                                                                       9 RAFVTIGK 16
                                                                     1 RAFVTIGK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neuroprotective
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(INRM ) INST
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13-APR-2000
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The present invention describes a composition capable of eliciting anti-
major histocompatibility (MHC) antibodies. The composition comprises an antigent associated with an altered MHC (lass II determinant (I) comprising alphal, alpha2, beta1 and beta2 polypeptide domains encoded by a mammalian MHC class II locus covalently linked to form a polypeptide (I) contraining beta2, alpha2, alpha1 and beta1 domains in sequence. The resulting Antigen-MHC complex is recognizable by the T cell receptor. The compositions are used for immunisation against, or treatment of, a wide compositions are used for immunisation against, or treatment of, a wide (AIDS), lupus erythematosus, multiple sclerosis, thyroiditis, toxic shock, tumour and snakebite, depending on the nature of antigen. (I) is also used to analyse functional interactions between the various domains and for targeting lymphocyte receptors. Antibodies against (I) are produced by usual methods of immunisation or cell fusion, and may be humanised by standard methods. These antibodies are useful for diagnosis (detection or purification of MHC gene produces), therapy (neutralising MHC on cell surfaces) and in the study of MHC and cellular processes.

AAZ33240 to AAZ33242 and AAY52840 to AAY52940 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Composition containing an antigen and altered major histocompatibility Class II determinant, used to immunize against autoimmune diseases, e.g. acquired immune deficiency syndrome.
                                                                                                                                                                                                                                                                                                                                 Major histocompatibility complex, MHC class I; MHC class II; antigen; immune response; diagnosis; antibody; immunisation; autoimmune disease; acquired immune deficiency syndrome; AlDS; cytostatic; dermatological; anti-inflammatory; neuroprotective; immunosuppressive; antithyroid; vaccine; lupus erythematosus; multiple sclerosis; thyroiditis;
                                                                                                                                                                                                                                                                                            Altered MHC determinant binding peptide SEQ ID NO:35.
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(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kourilsky P, Mottez E, Abastado J;
                                                                                                                                                                          AAY52857 standard; peptide; 16 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                      toxic shock; tumour; snakebite.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     91US-00792473
91US-00801818
                                                                                                                                                                                                                                                         (first entry
8; Conservative
                                                                            RAFVTIGK 16
                                       RAFVTIGK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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Matches
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                                                                                                                                                            AAY52857
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The present invention relates to a major histocompatibility complex (MHC) class I determinant, which has alpha_1 alpha_2 alpha_3 and beta2—microglobulin polypeptide domains encoded by a mammalian MHC class I locus. The MHC class I determinants are useful for activating the immune system and presenting antigens to the immune system to elicit an antigenic response. The MHC class I determinants are also useful for treating diseases, e.g. T cell mediated autoimmune disease, AlDS, lupus erythematosus, toxic shock or make bite. The altered MHC class I determinants and compositions containing antigens bound to the determinants are useful in diagnostic applications, e.g. altered determinants may be used to target lymphocyte receptors and the resulting bound determinant can be assayed. (Updated on 11-SEP-2003 to standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful for eliciting an immune response and/or for immunizing against or treating diseases, for example, multiple sclerosis, AIDS, toxic shock or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New altered major histocompatibility complex (MHC) class I determinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mottez E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                            Major histocompatibility complex; MHC class I; immune; snake bite; T cell mediated autoimmune disease; AIDS; lupus erythematosus;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 39; DB 4; Length 16; Pred. No. 0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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                                                                                                                                                                                                  Altered MHC determinant binding peptide #17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAT SANTE & RECH MEDICAL.
                                                                                                                                                                                                                                                                                          8
                                                                                                                                                                                                                                                                                       Human immunodeficiency virus; type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Col 11; 105pp; English.
                                                                                                AAB58618 standard; peptide; 16 AA
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91US-00801818.
93US-00072787.
93US-00117575.
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ilarity 100.0%;
Conservative 0;
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                                                                                                                                                                         (first entry)
                                                                                                                                                           (revised)
                           RAFVTIGK 16
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1 RAFVTIGK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 16 AA;
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(INRM ) INST
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13-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                           09-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                      15-NOV-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                     3-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUN-1993;
                                                                                                                                                                                                                                                             toxic shock
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Score 39; DB 3; Length 16; Pred. No. 0.14;

100.0%;

Query Match Best Local Similarity

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RESULT 85
AAY40414
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AAR42057
ID AAR42
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SXSXXXXXXXXXX
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Best Local Similarity
Matches 8; Conserv
Synthetic
                                                                                                                 AAY40414 standard; peptide;
                                                                                                                                                                                                                                                          Sequence 17 AA;
                                                                                                                                                                                                                                                                             Peptide P-18, opt. with a Cys-Gly dipeptide attached at its N- terminus, is the pref. peptide for use in raising a cytotoxic T lymphocyte response against HIV. The peptide is encapsulated in a liposome for admin. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 16; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                       Inducing cytotoxic T lymphocyte response to HIV - with liposome contg. peptide or protein having CTL epitope of HIV and protein, also improving humoral immunity, useful in vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-1993;
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                    Lipopeptide;
HIV related (
                                                                         25-NOV-1999
                                                                                              AAY40414;
                                                                                                                                                                                                                                                                                                                                                                                                                       Alving
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Immunodeficiency Virus type 1; envelope protein;
vaccine; AIDS; peptide P18; epitope.
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29-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR42057;
                                                   Lipopeptide
                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1993-336590/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                           (MEDI-) MEDIMMUNE INC. (USSA ) US DEPT ARMY.
                                                                                                                                                                       10 RAFVTIGK 17
                                                                                                                                                                                           1 RAFVTIGK 8
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                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                       Cassatt D,
                    antigen; cytotoxic T lymphocyte; steroid; vaccine; condition; tumor cell; NP protein.
                                                   comprising a fragment of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
(first en
                                                                         (first entry)
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3. .17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            label= P-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry
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0.15;
                                                  HIV env protein.
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                                                                                                                                                                                                                                    Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunogen;
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ID ADN14
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XX HIV h
XX HIV;
KW CTL;
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes lipopeptide that comprise a partial peptide containing 10-40 amino acids and at least one antigenic determinant specific for cytotoxic Tlymphocytes. The lipopeptide comprises at least one 10-20 carbon fatty acid derivatives and/or at least one modified steroid group. The lipopeptides are useful for: the preparation of a vaccine against HIV related conditions; immunizing a human or animal against an antigen by inducing cytotoxic T-lymphocytes; immunizing a human or animal against pathogens (especially a virus e.g. HIV-1 and HIV-2, or parasites). The present sequence represents a lipopeptide of the invention, and comprises part of the HIV env protein
                                                                                             HIV; antigen; epitope; T cell; MHC; major histocompatibility complex; CTL; cytotoxic T lymphocyte; HIV infection; cancer; tuberculosis; tumour; hepatitis; melanoma; breast cancer; Hodgkin lymphoma; nasopharyngeal carcinoma; vaccine; immune response; hyaluronic acid; HA; CD8+ T cell; viral infection; bacterial infection;
                           Human
                                                                                                                                                                                                                                                                HIV helper T cell epitope #42.
                                                                                                                                                                                                                                                                                                                    17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                        ADN14075;
                                                                                                                                                                                                                                                                                                                                                                                                                          ADN14075 standard; peptide; 17 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 4; Page 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New lipopeptide inducers of cytotoxic T lymphocytes, useful as against cancers, viruses, parasites and HIV-related conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-553128/47.
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18-DEC-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modified-site
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                           immunodeficiency virus
                                                                             infectiono;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RAFVTICK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90FR-00015870.
91EP-00403446.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "amidated
17
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                                                                             parasitic infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   35pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sergheraert C,
J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 39; DB 2;
Pred. No. 0.15;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Magne
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 æ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a C14
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RESULT 87
AAR31277
                                                                                                                                                                                                                                                                                                              considered therapeutic comprising administering to the mammal a particle comparison at least one peptide that comprises a Toell epitope comprises a sequence of at least about eight amino carid (MA) polymer analogue correcognised by a major histocompatibility complex molecule of the mammal. The Toell epitope comprises a sequence of at least about eight amino carids of the antigen. Also included are a method of improving major carids of the stocompatibility complex (MHC) presentation of a Toell epitope of an instruction in a mammal (comprising administering to the mammal the conjugate). The Toell epitope is recognised by a major can instruction or an HC Class I molecule and by a CD8+ Toell of the mammal. The immune system response comprises a cytotoxic T lymphocyte, a CD4+T cell, or an antibody that recognises the antigen. The immune system response to the antigen is an antigen of a pathogenic agent or a cumour cell. The immune system response to the antigen is an antigen of a pathogenic agent or a cumour cell. For modulating, i.e. enhancing or diministration and methods care useful for modulating, i.e. enhancing or diministration is also useful conjugates are useful as peptide vaccines against an antigen of a conjugates are useful as peptide vaccines against an antigen, a conjugates are useful as peptide vaccines against an antigen, a conjugates are useful as peptide vaccines against an antigen, a conjugates are useful as peptide vaccines associated with any of the cantigen above e.g. HIV infection, cancer, thus remains are carcinoma. The peptide vaccine compositions are canditional adjuvants, but still induce a stronger cell mediated response cumour cell mediated response compositions of the present invention do not require cadditional adjuvants, but still induce a stronger cell mediated response compositions of the present sequence is an HIV-1 and carcinome of the present sequence is an HIV-1 and carcinome of the present sequence is an HIV-1 and carcinome of the present sequence is an HIV-1 and carcinome of th
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                                                                                                                                                                                                                             Query Match
Best Local S
Matches 8
AAR31277;
                                   AAR31277 standard; peptide; 18 AA
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                                                                                                                                                                                                                                                                                                              Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modulating an immune system response to an antigen in a mammal, comprises administering a particle-free therapeutic comprising a hyaluronic acid polymer analogue covalently linked to a peptide that comprises a T cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Li FQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-AUG-2001; 2001US-0310498P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-FEB-2002; 2002US-00062710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-MAR-2003
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                                                                                                                                                                                   1 RAFVTIGK 8
                                                                                                                                                                                                                           Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chu Y,
                                                                                                                                            RAFVTICK 14
                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 12; 23pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     relates to modulating an immune system response to an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Qiu J;
                                                                                                                                                                                                                           100.0%; ;
100.0%; ;
htive 0;
                                                                                                                                                                                                                           Score 39; DB
Pred. No. 0.1
0; Mismatches
                                                                                                                                                                                                                                              0.15;
                                                                                                                                                                                                                                                                DB 7; Length 17
                                                                                                                                                                                                                           Indels
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RESULT 88

AAR30032 standard; peptide; 18 AA

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RAFVTIGK 8

BXAXB

25-MAR-2003 AAR30032;

(revised

Query Match
Best Local Similarity
Arches 8; Conservi

Conservative

0

Score 39; DB 2 Pred. No. 0.16; ; Mismatches

DB 2; 0,

Length 18

0,

Gaps

0

100.0%;

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The invention relates to a co-conjugate comprising an immunogenic protein cor protein complex having a first set of covalent linkages to low molecular weight moieties which have an anionic or polyanionic character at physiological pH, and a second set of covalent linkages to peptides comprising HIV principal neutralizing determinants (PND's) or immunologically equivalent peptides. Preferably at least one set of the covalent linkages is comprised of maleimide derivatives; the polyanionic moiety is composed of one to five residues of the anionic form of a carboxylic, sulphonic or phosphonic acid, the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b; and the PND peptide has a linear structure, a disulphide-bonded cyclic structure. The present sequence (PND135-18) is an example of a PND peptide component used in the co-conjugate. The co-conjugate is useful for inducing anti-peptide immune response in mammals, for formulating vaccines to prevent HIV infection or disease, including AIDS, or for treating humans afflicted with HIV infection or disease.
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19-JUL-1990;
19-JUN-1991;
19-JUN-1991;
Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 12; Page 56; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conjugate proteins comprising HIV peptide components - useful for preparing vaccines for e.g. AIDS or for treating infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1992-026505/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Leanza WJ, Marburg S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-JUL-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP467700-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV principal determinant peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MERI ) MERCK & CO INC.
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ΑA;
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90US-00555966.
91US-00715276.
91US-00715278.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tolman RL,
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RESULT 89
AAR26713
ID AAR267
XX AR26
XX O9-FB
XX HIV-P
XX Human
XW                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                          HIV-PND-polysaccharide-protein conjugate vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The peptide is HIV principle neutralising determinant (PND) 135-18 and is used as part of a conjugate comprising the major immune enhancing protein (MIEP) of Neisseria meningitidis covalently linked to the HIV PND. The conjugate may be used to prepare vaccines against HIV infections, e.g. AIDS, as research tools for studying PND structure- function relationships, or as immunogens for use in the passive immunisation of humans. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conjugates of HIV-1 PND peptide(s) meningitidis - useful as a vaccine infection, e.g. AIDS in humans.
                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                 outer
                                                                                                                                                                                                                                                                                                                                            Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                09-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR26713;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR26713 standard; peptide; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 AA;
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                             29-JAN-1992
                                                                              EP468714-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1992-425771/52.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-JUN-1992;
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                                                                                                                                                                                  Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-APR-1993
                                                                                                                                                                                                                                                                                                           immunodeficiency virus; principal neutralizing determinant; membrane protein complex; OMPC; Neisseria; AIDS; PND-135-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9; Page 59; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immune enhancing protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunodeficiency
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 100
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MERCK & CO INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91US-00715273
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                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                            /note= "Joins onto polysaccharide-protein complex via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Marburg
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                                                                                                                                 gite"
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protein; vaccine; anti-HIV antibodies; immunogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tolman RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               with the MIEP of Neisseria for treating and preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.16;
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Matches
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                                                                                                                                           determinant (PND), or an immunologically equivalent peptide (PEP), covalently coupled to an immunogenic protein or protein complex through an anionic polysaccharide linker. Pref. the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b and the PND peptide has a linear structure, a disulphide-bonded cyclic structure, an amide-bonded cyclic structure, a thiosther-bonded cyclic structure. The present sequence (PND135-18) is an example of a PND peptide component. The conjugates are used for inducing HIV-neutralising antibodies or for making vaccines to prevent contraction of HIV infection or disease. The antibodies can be used for passively protecting against infection, or for treating AIDS, or in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUL-1990;
19-JUL-1990;
19-JUN-1991;
19-JUN-1991;
                                                                                                                    Sequence 18
                                                                                                                                                                                                                                                                                                                  The invention relates to a conjugate of an HIV principal neutralizing determinant (PND), or an immunologically equivalent peptide (PEP),
                                                                                                                                                                                                                                                                                                                                                                                        HIV peptide-polysaccharide-protein conjugates - used in vaccines produce antibodies to prevent or treat HIV infection.
                                                                                                                                                                                                                                                                                                                                                            Claim 9; Page 57; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MERI ) MERCK & CO INC.
                                                                         Local
                                                      l Similarity
8; Conserva
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RAFVTICK 15
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90US-00555974.
91US-00715275.
91US-00715277.
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                                                                      Score 39;
Pred. No.
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ARESULT 90
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DT 24-OC
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DT 20-MA
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AR1tig
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KW Itpop
KW Cytot
KW Pathc
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Humar
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PR 03-MJ
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PR 01-MJ
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PR 01-MJ
XX
PR (UYR) 24-OCT-2003 25-MAR-2003 20-MAY-1994 AAR44190; AAR44190 standard; peptide; 18 (revised) 8

0,

Antigen; B2; third variable domain; V3 loop; gp120; HIV-1; vaccine; strain IIIB; multiple antigenic peptide system; dendritic core; lipophilic membrane anchoring group; mammal; humoral; immunisation; cytotoxic T cell; CT; immune response; infection; Freunds adjuvant; pathogen; HIV; influenza; malaria.

gp120 V3 loop

(first entry) antigen B2.

Human immunodeficiency virus 1.

11-NOV-1993 93WO-US004179

01-MAY-1992;

92US-00877613.

(UYRQ ) UNIV ROCKEFELLER

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RESULT 91
AAR58548
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                        16-OCT-2003
25-MAR-2003
29-MAR-1995
                                     New recombinant proteins contg multiple antigenic determinants - by flexible hinge domains.
                                                                                     WPI; 1994-279687/34.
                                                                                                                                                                               10-FEB-1993;
                                                                                                                                                                                                            10-FEB-1994;
                                                                                                                                                                                                                                          18-AUG-1994.
                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                  PKK-MEAV
                                                                                                                                                                                                                                                                                                                                            HIV-1; V3 loop; multiple epitope; AIDS; vaccine; MEAV; Escherichia coli;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR58548;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New multiple antigen system esp. for use in HIV vaccines - contains lipophilic membrane anchor imparting adjuvant activity, and peptide antigens coupled to dendritic core.
                                                                                                                                                 (UNBI-) UNITED BIOMEDICAL INC
                                                                                                                                                                                                                                                                        WO9418234-A1
                                                                                                                                                                                                                                                                                                                                                                            HIV-1 isolate IIIB V3 loop domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; Page 27; 55pp; English.
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(revised)
(first entry)
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Pred. No. 0.16;
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Disclosure; Page 36; 56pp; English

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RESULT 92
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ID ANM63
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Best Local S
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                                                    cancer which comprises incubating lymphocytes with a tumour-specific antigen or autoimmune disease related antigen and a superantigen. The treated cells are then introduced into the patient. The superantigen and the tumour-specific antigen or autoimmune disease related antigen can be conjugated together. The products are used to treat cancer (carcinoma, melanoma, lymphoma etc.), infections (bacterial, viral or eukaryotic) and autoimmune disease (e.g. idiopathic thrombocytopaenic purpura, rheumatoid arthritis, systemic lupus erythematosus, multiple sclerosis etc.). The antigens either induce an immune response or inhibit a patholgical
                                                                                                                                                                                                                                                                                                                                          Synthetic peptides AAW63049-85 are used, with superantigens, to exthe invention. The specification describes a method for treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conjugates and polymers containing superantigen and therapeutic antigen for treatment of cancer, infection, autoimmune disease and graft rejection, also treatment by administering lymphocytes treated in vitro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEAV gene (AAQ70535) encodes a portion of the CD4 binding domain (AAR58550) of HIV env protein, the domain being capable of inducing a helper T- cell response, and 5 peptide domains from the V3 loop of HIV-1 isolates MN, SC, RF, IIIB and MMJ2 (AAR58545-49), each peptide being separated by a spacer domain (AAR58551). The gene was expressed in E. coli BL21/pKK-MEAV for preparation of a multiple epitope AIDS vaccine (AAR58552). (Updated on 25-MAR-2003 to correct PN field.) (Updated on 16-
                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Page 40; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-362497/31.
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17-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       autoimmune disease related antigen;
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8; Conserv
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97US-0044074P.
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Pred. No.
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inhibit; pathological response;
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                                                                                                                                                                                                                                                                                                                                                                       to exemplify
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RESULT 93
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Best Local
                                                                                                   The present sequence comprises a glycosylated fragment of gp120. Glycopeptides comprising a fragment of the MUC1 repeat sequence, especially having a Gal-GalNAc or GalNAc moiety on Thr-10 or Thr-17 (see AAY96172-74), are useful as immunomodulators, causing super-proliferation of T cells. Such glycopeptides can be used in the treatment or diagnosis of a disease, in particular cancer, or as vaccine adjuvants. The glycopeptides are particularly useful in manufacturing a medicament for preventing or treating cancer by stimulating T cells whose receptors recognize the glycopeptide. They are also useful for diagnosing or treating cancer, e.g. carcinoma (e.g. mammary, lung, bladder or colon carcinomas, or ovary and endometrial tumours), or sarcoma (e.g. soft tissue and bone sarcomas, or leukaemia). Human peripheral blood lymphocytes were used in a proliferation assay. The proliferation index of the gp120 glycopeptide (taking the index as 1 when no glycopeptide was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18
                                                                                                                                                                                                                                                                                                                                                                   New immunomodulating glycopeptide that causes super-proliferation of T cells, useful for treating cells in vitro, for diagnosing or treating cancer (e.g. carcinoma or sarcoma) or as an adjuvant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Burchell J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200052046-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gp120; MUC1; immunomodulator; glycopeptide; T-lymphocyte; T-cell;
proliferation; cancer; sarcoma; carcinoma; leukaemia; diagnosis; therapy;
vaccine; adjuvant; glycosylation.
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                                                             Sequence 18
                                                                                         of the gp120 glycopeptide (taking the present) was 1-1.7
                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 24; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-601868/57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAR-2000; 2000WO-GB000724
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Local Similarity
les 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Taylor-Papadimitriou J;
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gp120 glycosylated peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99GB-00004695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "O-glycosylated by GalNAc-beta-1-3Gal"
                100.0%; Score 39; DB 3
100.0%; Pred. No. 0.16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 39; DB 2
100.0%; Pred. No. 0.16;
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                              DB 3;
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                              Length 18;
   Indels
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ARBSIJ13
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CX IR-DE
CX IR-DE
CX IR-DE
CX Vacci
PT HIV,
PT CYCOT
CC INCL
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                                                             Matches
                                                                                    Query Match
Best Local (
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18-DEC-1991;
18-DEC-1991;
                                                                                                                                                                                                                                                  The present sequence is a lipopeptide, which can be used for the immunisation of humans or animals against tumour cells or pathogens, specifically viruses, especially HIV-1 or HIV-2. The pathogens may also include parasites. Examples illustrate immunisation of mice against influenza, as well as HIV. The lipopeptide, with the appropriate antigenic determinants, can induce a strong cytotoxic T-lymphocyte response in a host organism against a wide range of pathogens. Addition
                                                                                                                                                                                                                          response in a host organism against a of an adjuvant is unnecessary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vaccine for immunization against tumor cells or pathogens, especially HIV, comprising peptide part, antigenic determinant specifically inducing cytotoxic T-lymphocytes and N-palmitoyl-lysine-derived chain(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Boutillon C, Martin
Gomard E, Tartar A,
                                                                                                                                                                      Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Page 17; 31pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-114040/13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EP1065212-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lipopeptide; cytostatic; virucide; anti-HIV; antiparasitic; vaccine; immunisation; tumour; pathogen; virus; antiviral.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lipopeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modified-site
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(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
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     1 RAFVTIGK 8
                                                       8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RAFVTIGK 8
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                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            #2 used in a vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Martinon F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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91EP-00403446.
99EP-00105773.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /label= Xaa
/note= "Xaa is optionally 2-acetylamino-hexadecanoyl,
/note= "Xaa is optionally 2-acetylamino-hexadecanoyl,
-bis(hexadecanoylamino)butyryl, or not present"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                    100.0%; Score 39; 100.0%; Pred. No.
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J;
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                                                             Mismatches
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0.16;
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                                                                                                              Length 18;
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RAFVTIGK 17

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RESULT 95
AAW24218
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                                                                                                                                                                                                          CC Peptide fragments AAW24217-20 were used to demonstrate a new method of CC tracing sources of infectious diseases. The method comprises preparing a CC short (9-50 amino acid) peptide containing at least one non-conserved CC epitope of an organism, injecting a composition containing the peptide CC intradermally into a test subject in a delayed-type hypersensitivity CC (DTH) assay and observing the injection site at intervals for induration. CC In this example CO4+T-lymphocyte epitopes to the HIV-1 VJ loop were CC mapped by existing in vitro technique for two existing HIV infected individuals and used to design peptides AAW24217-20. The method allows CC the T-lymphocyte epitopes of a large antigen to be determined in vivo in CC numans. The method is useful in medicine e.g. in diagnosis, monitoring CC and treatment design for infectious disease exposure, active autoimmune CC disease, allergic diseases and malignancy. It is especially useful for tracing infectious diseases and malignancy. It is especially useful for CC racing infectious diseases e.g. HIV, particularly when a sequence is present only in certain strains of an organism, and developing suitable CC vaccines. Vaccinated individuals can also be tested to verify protection CC against a particular strain. The method allows in vivo mapping of T-lymphocyte epitopes, not previously possible. The method is simpler, more capting and more sensitive. It can also be applied in a variety of inconversers of undarelegate contents.
                                                                                    Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide fragments containing antigen epitope(s) used to trace diseases used in a delayed-type hypersensitivity assay, for in vivo mapping of human T-lymphocyte epitope(s) e.g. for diagnosis, vaccine development
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
Region
                                                                                                                                           Sequence 19 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T-lymphocyte
delayed-type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 6; 14pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-393814/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (USSA ) US DEPT ARMY GOVERNMENT US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-JAN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD4+ T-lymphocyte epitope to HIV-1 V3 loop derived peptide V3-LAI-B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-MAR-1998
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4
                                1 RAFVTICK 8
                                                                    Similarity 100
8; Conservative
RAFVTIGK 11
                                                                                                                                                                                               e.g. undeveloped regions since specialist equipment is not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   epitope; diagnosis; antigen; infectious disease; hypersensitivity assay; vaccine development.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                    100.0%;
                                                                    0;
                                                                                      Score 39;
Pred. No.
                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ARMY MEDICAL
                                                                                      0.17;
                                                                                                     DB 2;
                                                                    0
                                                                                                     Length 19;
                                                                    Indels
                                                                  0,
                                                                  Gaps
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AAR60203 ID AAR XX

AAR60203 standard; protein;

20

RESULT 97

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Query Match Best Local S Matches 8

Local Similarity

8

Conservative

0;

Mismatches

<u>,,</u>

Gaps

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100.0%;

100.0%; Score 39; DB 2; 100.0%; Pred. No. 0.18;

Length 20 Indels

1 RAFVTIGK 8

RAFVTIGK 10

Sequence 20

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RESULT 96
AARO4434
AC AARO444A
AC AARO4
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Lynn DU, P
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01-JUN-1989;
19-SEP-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-SEP-2004
25-MAR-2003
20-SEP-1990
                                                                                                                                                         Peptide 132 comprises segments of the Principal Neutralising Domain (envelope protein) from isolate HIV-IIIB. A Cysteine can be added, so that the residues at or near both ends can form a disulfide bond, thus giving the peptides a loop configuration. The loop configuration can be utilized to enhance the immunogenic properties of the peptides. The protein is capable of eliciting, and/or binding with, neutralising antibodies. The neutralising domain is bounded by cysteine residues which occur at positions 296 and 331. The peptides can be used as immunogens or screening reagents to generate or identify poly- or monoclonal antibodies. The last Tyr residue is an immunologically silent spacer. See also ARR0427-R04506 and ARQ04273-Q04279. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.) (Updated
                                                                   Revised record issued on 09-SEP-2004 :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Principal neutralising domain of HIV variants - used for producing peptide(s) and antibodies for diagnosis, prophylaxis and/or therapy therapy of HIV infection.
                                                                                                                                       on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8 (37); Page 76; 108pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-OCT-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV-IIIB; peptide 132; principal neutralising domain; antibodies; diagnosis; prophylaxis; therapy; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR04434;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-APR-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (REPK ) REPLIGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Petrobre J;
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(revised)
(first entry)
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89US-00359543.
89US-00407663.
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                                                                       Correction
                                                                   to Feature Table Key
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ID AAW76
XX AAW76
AC AAW76
AC AAW76
XX 25-JA
XX 25-JA
XX E Cel
KW bumar
KW proph
KW proph
KW micrc
KW antiv
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Best Local S
Matches 8
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25-MAR-2003
13-MAR-1995
               B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                     A molecular tag was created by annealing two complementary 76mer oligonucleotides with cohesive end overhangs. AAQ70167 is the sense strand and includes a BclI overhang, the HV gpl10 V3 loop coding sequence and a stop codon. The peptide encoded by the molecular tag (AAR60203), when part of a single chain fusion protein with binding regions from different antibodies, affected the avidity and binding specificity of the antibodies. For example, the tag failed to function properly when fused to L6 but performed successfully when fused to CD3. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG-2003 to
                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Expression vector encoding bispecific fusion protein - having binding domains for separate targets joined by helical peptide, useful e.g. for diagnosis and treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-FEB-1993;
13-SEP-1993;
                                                                                                   Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #83.
                                                                                                                                                            AAW76943;
                                                                                                                                                                                       AAW76943 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-250885/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  fusion protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV gp110 V3
                                                                                                                               25-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR60203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ledbetter JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BRIM ) BRISTOL-MYERS SQUIBB CO.
                                                                                                                                                                                                                                                                                                                  Local Similarity
nes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   n protein; recombinant bispecific single chain antibody; immunodeficiency virus; glycoprotein gp110; V3 loop.
                                                                                                                                                                                                                                                               12
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                                                                                                                                                                                                                                                             RAFVTIGK 19
                                                                                                                                                                                                                                                                                                                                                                                                          field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 12; 50pp; English.
                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             loop molecular tag.
                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gilliland LK, Hayden MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93US-00013420.
93US-00121054.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94EP-00300692
                                                                                                                                                                                                                                                                                                                                  100.0%; Score 39; 100.0%; Pred. No.
                                                                                                                                                                                        20
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RESULT 99
AAW54930
ID AAW54
XX AAW54
AC AAW54
XX Z5-SE
XX ITMU
DE HIV 9
XX ITMU
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PD 09-JI
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PR 09-JI
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Best Local S
Matches 8
 12-FEB-1988;
01-MAR-1991;
09-JUL-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, so prevents or inhibits 'bystander' apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral
                                                            07-JUN-1995;
                                                                                         09-JUN-1998
                                                                                                                                                  Human immunodeficiency virus
                                                                                                                                                                              Immunoadsorbent; immunoassay;
                                                                                                                                                                                                           HIV gp120 envelope protein, peptide 127, analogue 127h'.
                                                                                                                                                                                                                                        25-SEP-1998
                                                                                                                                                                                                                                                                      AAW54930;
                                                                                                                                                                                                                                                                                                AAW54930 standard; peptide; 20 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T cell immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 39; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce telerance to the epitopes for treatment of human immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-506315/43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-FEB-1997;
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Homo sapiens
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8; Conserv
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                                                                                                                                                                                                                                        (first entry)
   88US-00155321.
91US-00663262.
91US-00726605.
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                                                            95US-00488252
                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 39;
100.0%; Pred. No.
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                                                                                                                                                                               VIH
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                                                                                                                                                                              gp120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.18;
                                                                                                                                                                            immunogen; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2;
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Best Local
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                                                                                                                                                                                                                                                                                                                   three-dimensional atomic structural conformation; protein co-ordinate data; V3 loop peptide; HIV-1; envelope glycoprotein; gp120; human monoclonal antibody 447-52D; murine monoclonal antibody 0.5 beta; immunogen; immunogenic; V3 loop epitope; HIV-1 infectivity inhibitor; anti-HIV; vaccine;
                                                            Composition for inhibiting HIV-1 infection, comprises isolated peptide molecule that mimics atomic structural conformation of V3 loop peptide HIV-1 envelope glycoprotein that is bound to, and constrained by human monoclonal antibody.
                                        Example 1;
                                                                                                                 WPI; 2004-625447/60.
                                                                                                                                     Anglister J,
                                                                                                                                                                                           04-FEB-2003; 2003US-0444682F
                                                                                                                                                                                                              04-FEB-2004; 2004WO-US003304
                                                                                                                                                                                                                                                          WO2004069863-A2
                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                      HIV-1 V3-IIIB related peptide SEQ ID NO:37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADR18886 standard; peptide; 20 AA
                                                                                                                                                                                                                                     19-AUG-2004.
                                                                                                                                                                                                                                                                                                             HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                             04-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                  ADR18886;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptides AAW54903-W54941 can be used as immunoassay for detecting antibodies to eliciting antibodies to HIV in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV gp120 peptides - useful as immunoassay reagents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                            UNIV NEW
YEDA RES
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8; Conserv
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                                        SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                    Sharon M, Schapira M,
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                                                                                                                                                                     YORK STATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21/22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Sc.
100.0%; Pr
                                       37;
                                                                                                                                                           CO LTD
                                     127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 39; DB 2;
Pred. No. 0.18;
0; Mismatches
                                                                                                                                    Zolla-Pazner S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           an immunoadsorbent in an HIV gp120, or as an immu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>ب</u>
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                                                                                                                                     Rosen
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The present invention describes a composition (C1) which comprises an isolated peptide molecule or isostere that mimics the three-dimensional (3D) atomic structural conformation of the V3 loop peptide of the HIV-1

(CONN-) CONNAUGHT LAB LTD

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RESULT 101
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                                                                15-AUG-1994;
                                                                                                                             15-AUG-1995;
                                                                                                                                                                                                                                                                     WO9605292-A1
                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR93073;
                                                                                                                                                                                                        22-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                       Antigen; non-infectious; retrovirus; long terminal repeat; gag; pol; env;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antigenic peptide CLTB73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR93073 standard; peptide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in an infected subject. The the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          binding fragment of the MAD, where the constrained V3 loop peptide differs in conformation from the same V3 loop peptide when it is in free form. Also described: (1) identifying (M1) from several existing compounds a molecule that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 correceptor on the surface of a receptor-bearing target cell; (2) designing a molecule that is useful as an HIV-1 V3 loop immunogen or as an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          envelope glycoprotein gp120 that is bound to, and const
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8; Conserv
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                                                                94US-00290105
                                                                                                                                    95WO-CA000483.
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AIDS; HIV; antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       immune response;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                     UDP-N-acetyl-alpha-D-galactosamine;
polypeptide N-acetylgalactosaminyltransferase; GalNAc-t3; human;
glycosylation; HIV-V3.
                                                                                                                                                                                                                                                                                                                                                                                       11-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 AA;
 WPI; 1998-008874/01
                               Clausen H,
                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antigenically marked non-infectious retrovirus-like particles - used to vaccinate against, and in the treatment of, AIDS and AIDS related
                                                                                                              15-MAY-1996;
                                                                                                                                                                                                           WO9743405-A1
                                                                                                                                                                                                                                         Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW34475 standard; peptide; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             used to identify antibodies specifically reacting with retrovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 4; Page 38; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rovinski B,
                                                                                                                                            15-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                    Acceptor peptide HIV-V3.
                                                               (CLAU/) CLAUSEN (BENN/) BENNETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |||||||
RAFVTIGK 21
                               Bennett EP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                     (first entry
                                                                                                              96US-00648298
                                                                                                                                             97WO-DK000226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Persson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 39; DB 2
Pred. No. 0.19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ά,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Klein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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New isolated N-acetyl-galactosaminyl-transferase enzyme - used for the production of glycosylated polypeptide(s) having particular enzymatic, immunogenic or other biological or physical properties. used for the

Example 2; Page 30; 70pp; English.

Acceptor peptides Muc2, Muc5c (see AAW34474) and HIV-V3 (see AAW34475) were used to study the acceptor substrate specificity of the novel human N-acetylgalactosaminyltransferase GalNAc-T3 (see AAW34470). Expression of a soluble GalNAc-T3 construct in Sf9 cells resulted in significant increases in GalNAc-T3 construct in Sf9 cells resulted in significant infected cells compared to uninfected controls or cells infected with the host-blood group O2 gene. GalNAc-transferase activity with the Muc2 acceptor peptide was increased 20-fold, and activity with the HIV-V3 peptide was increased nearly 100-fold. In contrast, expression of GalNAc-T1 and -T2 constructs only increased the GalNAc-transferase activity coward Muc2 and Muc5C peptide substrates. This illustrates the unique acceptor substrate specificity of GalNAc-T3. The enzyme is used in claimed methods for the glycosylation of peptides and proteins and for producing vaccines by modifying the O-glycosylation pattern of eukaryotic

Sequence 21 AA;

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Matches
                                                 Query Match
                                        Local
10
            1 RAFVTIGK 8
                                 Similarity
8; Conserv
RAFVTICK 17
                                  Conservative
                                        100.0%; Score 39; 100.0%; Pred. No.
                                  0;
                                   Mismatches
                                         0.19;
                                                  DB 2;
                                  <u>,</u>
                                                  Length 21;
                                  Indels
                                  <u>.</u>.
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S

AAW79180 standard; peptide; 21

25-JAN-1999 (first entry)

Fusion immunoglobulin heavy chain HIV gp120 T cell epitope #58.

RESULT 103
AAW79180
ID AAW791
XX AAW791
XX AAW791
XX Pusion
XX B cell
XX B cell
XX Muman
XX microb
XX lance
PN 13-FEE
XX lance
PN lance B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.

Homo sapiens

WO9836087-A1.

20-AUG-1998

13-FEB-1998; 98WO-US002766

13-FEB-1997; 97US-0040581P.

(AMNA-) AMERICAN NAT RED CROSS

Scott D, Zambidis E;

WPI; 1998-506315/43

New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency

Disclosure; Page 50; 154pp; English

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RESULT 104
AAW76901
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
                                                                                                                                                                                                                                                                               New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, unduce tolerance to the epitopes for treatment of human immune defined to the contract of the section of the sect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, so prevents or inhibits 'bystander' apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-506315/43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AMNA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9836087-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fusion immunoglobulin heavy chain HIV gp120 T cell epitope #20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW76901;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW76901 standard; peptide; 21 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , sp.
no sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AMERICAN NAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RAFVTIGK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97US-0040581P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98WO-US002766.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RED CROSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 39; DB 2
Pred. No. 0.19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                             used to
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This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (19H) protein with a mammalian, especially human, 19H chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) ppl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection,

ç

Claim

154pp; English

deficiency

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RESULT 105
RAW75478
ID AAW754
XX AAW754
XX AW754
XX IT-OCT
DT 17-OCT
DT 20-MAR
DT 27-APR
DT 27-APR
XX V3 loc
KW V3 loc
KW HTLV-I
XX HUMAN
PR 15-JUN
PR 15-JUN
PR 09-JUN
XX IF INMUNC
PR 15-JUN
PR 09-JUN
XX IF INMUNC
PR 15-JUN
PR 15-JUN
PR 15-JUN
PR 09-JUN
XX IF INMUNC
PR 15-JUN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
                                                            self-assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle. The retrovirus-like particle comprises a modified HIV genome devoid of long terminal repeats (LTRs) and contains a nucleotide sequence coding for a chimeric envelope glycoprotein. The chimeric envelope glycoprotein has the HIV-1 gp120 conserved region 2 and a second retroviral envelope amino acid sequence from a heterologous strain of HIV-1, HIV-2, HTLV-I or HTLV-II inserted into the first retrovirus-like particle is useful in vaccines against HIV. (Updated on 20-MAR-2003 to correct PA field.) (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-OCT-2003
20-MAR-2003
27-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence represents a peptide from the V3 loop of the gp120 protein from the human immunodeficiency virus type 1 (HIV-1) strain HXB2. The peptide is used to determine antibody responses after immunisation with a peptide is the complex of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; Col 7-8; 12pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunogenic retrovirus-like particle - with chimeric HIV-1 envelope protein containing heterologous retroviral amino acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUN-1992;
09-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-FEB-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         V3 loop; gp120 protein; HIV long terminal repeat; LTR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1 strain HXB2 gp120 V3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW75478;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HTLV-II; vaccine; human T-lymphotropic virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CONN-) CONNAUGHT LAB LTD.
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RAFVTIGK
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(revised)
(first en
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93US-00073526.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; HIV-1; retrovirus-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95US-00453745.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  œ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Haynes J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chimeric;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           loop peptide amino acids 302-322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ..
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rovinski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         envelope;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     particle; genome; HIV-2;
glycoprotein; HTLV-I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                             retrovirus-
                                                                                                                                                                                                                                  l a second
in of HIV
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RESULT 106
AAY16052
ID AAY160
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Best Local Similarity
""" hes 8; Conserve
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                      S
                                                               Query Match
Best Local Similarity
                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-OCT-2003
20-MAR-2003
04-AUG-1999
                                                                                                                              The specification describes a nucleic acid molecule encoding a self assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle. The retroviral particle comprises a modified HIV genome devoid of long terminal repeate containing a nucleotide sequence coding for a chimeric envelope glycoprotein which has a first (a) and second (b) retroviral envelope amino acid sequence, where (a) contains the HIV-1 gpl20 conserved region 2, and (b) contains a retroviral envelope amino acid sequence of a heterologous strain of HIV-1, HIV-2, HILV-1 or HILV-II inserted into (a) at an endogenous site (BglII and StuI). (b) may comprise peptides AAY16049-51 and AAY16055. The nucleic acids are useful as vaccines against HIV. The present sequence is used in the course of the invention. (Updated on 20-MAR-2003 to correct PR field.) (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Retrovirus-like particle; modified HIV genome; chimeric envelope glycoprotein; HIV-1 gpl20; conserved region 2; HIV-1; HIV-2; HTLV-I; HTLV-II; vaccine.
                                                                                                                                                                                                                                                                                                                                                      Immunogenic retrovirus like particles comprising modified HIV genomes, useful as vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUN-1992;
09-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY16052 standard; peptide; 21 AA.
                                                                                                          Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                            Example 4; Col 9-10;
                                                                                                                                                                                                                                                                                                                                                                                                                          Cao SX,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US5912338-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV-1 isolate HXB2 gp120 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY16052;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ROVI/) ROVINSKI B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14 RAFVTICK 21
14 RAFVTIGK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 RAFVTIGK 8
                                                    80
                        RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                                                                                          Klein MH,
                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92US-00839751.
93US-00073526.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                          Наупев J,
                                                                                                                                                                                                                                                                                                                            12pp; English
                                                    100.0%; Score 39; DB 2
100.0%; Pred. No. 0.19;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 39; DB 2
Pred. No. 0.19;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                           Rovinski
                                                                              DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>.</u>
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                                                                              Length 21;
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                                                       Indels
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                                                    Gaps
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RESULT 107
AAW85568
ID AAW855
XX AAW855
XX 20-MAR
DT 20-MAR
DT 24-FEB
XX IMMUNO
KW IMMUNO
KW Chimer
KW Chimer
KW Chimer
KW HTLV-1
XX 30-MAY
XX 10-MAY
XX 10-
                                                                                                                                                                                                                                         RESULT 108
AAB15012
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents a Human immunodeficiency virus type 1 derived peptide. The peptide is used in the immunoassay diagnostic kit of the invention. The specification describes an immunoassay diagnostic kit for detecting antibodies in a sample, which comprises an antigen consisting of a self-assembled, non-infectious, non-replicating, immunogenic, retrovirus-like particle encoded by a modified HIV genome that is devoid of long terminal repeats and contains a nucleotide sequence containing HIV-1 gp120 conserved region 2 and a second amino acid sequence containing an envelope sequence of a heterologous strain of HIV-1, HIV-2, HTUV-I or HTUV-II. (Updated on 20-MAR-2003 to correct PR field.) (Updated on 20-MAR-2003 to correct PR
HIV; immune; diphosphonate
                                                                                                          07-DEC-2000
                                                                                                                                                                AAB15012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus type 1 derived peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-MAR-2003
24-FEB-1999
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                                                  Peptide P18 derived from V3
                                                                                                                                                                                                               AAB15012 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-069713/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUN-1992;
09-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US5849475-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HTLV-I; HTLV-II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; Col 9-10; 12pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    retrovirus-like particles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunoassay diagnostic kit for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                      14
                                                                                                                                                                                                                                                                                                                                                                                        1 RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                      RAFVTICK 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cao SX,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
ilarity 100.0%;
Conservative 0;
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(first entry)
                                                                                                          (first entry)
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93US-00073526.
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                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                     loop of HIV IIIB group 120 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 39; DB 2
Pred. No. 0.19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              detecting antibodies - comprising chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               $
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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RESULT 109
AAU08699
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Best Local
                       WPI; 2001-595518/67.
Differentiating between infection by human
                                            Rovinski B,
                                                                                      23-FEB-1998;
                                                                                                           23-FEB-1998;
                                                                                                                                18-SEP-2001.
                                                                                                                                                     US6291157-B1
                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                     Human immunodeficiency virus; HIV; retroviral antigen; gag; pol; immune response; antigenic marker; antigenic epitope; retrovirus
                                                                                                                                                                                                                                                              18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                immune response to an antigen. The present sequence is an immunoge peptide used to test the ability of the compounds to cause an income the immune response. The peptide consists of an amino terminal cycresidue, a glycine/alanine/glycine spacer and amino acids 308-322 V3 loop of HIV IIIB gp120 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The
                                                                                                                                                                                   Human immunodeficiency virus
                                                                                                                                                                                                                                                                                  AAU08699
                                                                                                                                                                                                                                                                                                      AAU08699 standard; peptide; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21
                                                                 (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to diphosphonate compounds useful as immunological adjuvants. The compounds can be used for stimulating an immune response to an antigen. The present sequence is an immunogeneic in the compounds to cause an increase in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         stimulating an immune response to an antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New diphosphonate compounds, useful as immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-514809/46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hawkins LD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EISA ) EISAI CO
                                                                                                                                                                                                                                                                                                                                                           13
                                                                                                                                                                                                                                                                                                                                                                                                     8
                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8; Page 86; 130pp; English.
                                                                                                                                                                                                                                                                                                                                                                              RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                           RAFVTICK 20
                                                                                                                                                                                                                                                                                                                                                                                                                                              Ä
                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                          Cao S,
                                                                                                                                                                                                                                                            (first entry)
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                                                                                     98US-00027955
                                                                                                           98US-00027955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0118131P
                                                                                                                                                                                                                                     particle CLTB73 containing a V3 (HXB2) antigenic marker
                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                            Yao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ST,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      virus.
                                            'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lewis M,
                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                          Persson
                                                                                                                                                                                                                                                                                                                                                                                                   Score 39; DB 3
Pred. No. 0.19;
; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcguiness
                                            Klein
                                                                                                                                                                                                                                                                                                                                                                                                                       DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nault
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   adjuvants
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cysteine
                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                env;
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                                                                                                                                                                                                                                                                                                                                                                                                   0
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immunodeficiency virus

Antibodies able to neutralise more The gp120 V3 loop sequences from di Example 16; Page 100; 154pp; English

se more than one HIV-1 isolate are claimed. from different isolates comprising the

New recombinant human antibody - with HIV neutralising activity against at least two isolates, useful for preventing or treating infection in

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-OCT-2003
25-MAR-2003
27-APR-1994
                                                                                                                                                                                                                    01-APR-1992;
                                                                                                                                                                                                                                                                                                                                                       Human Immunodeficiency Virus; antigen; ELISA; recombinant antibody; HIV-neutralising monoclonal antibody; immunoglobulin; AIDS; acquired immune deficiency syndrome; chimeric antibody; surface glycoprotein gp120; V3 loop; epitope mapping.
                                                                                                                              WPI; 1993-336600/42.
                                                                                                                                                           Emini EA,
                                                                                                                                                                                                                                            23-MAR-1993;
                                                                                                                                                                                                                                                                          14-OCT-1993
                                                                                                                                                                                                                                                                                                       WO9319785-A1.
                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                 gp120 V3 loop sequence of HIV-1 IIIB isolate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR42153;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR42153 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               generated by immunisation with an immunogenic composition capable of eliciting HIV-specific immune response. The antigenic marker may comprise at least one antigenic epitope from another virus. This sequence represents a retrovirus-like particle containing an antigenic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       specific antibodies with a non-infectious, non-replicating, immunogenic HIV-like particle as an antigen. The antigen comprises an assembly of a gag gene product, a pol gene product and a modified env gene product containing a non-retroviral heterologous, antigenic, anchor sequence that replaces the endogenous anchoring functions of the very gene product. The method detects immune complex formation between HIV-specific antibodies and the antigens. The method is also useful for identifying antiserum
                                                                                                                                                                                       (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibodies specifically reactive with HIV retroviral antigens in a sample. This involves contacting a sample suspected of containing HIV-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and antiserum generated by immunization against HIV, comprises use of infectious, non-replicating HIV-like particle with heterologous,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method for determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Col 17; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antigenic anchor sequence.
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                                                                                                                                                         Conley AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
                                                                                                                                                                                                                    92US-00861701
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                                                                                                                                                         Mark GE,
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                (IIIB isolate)
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                                                                                                                                                         Johnson LS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.19;
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                                                                                                                                                         Pfarr DS
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RESULT 111
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DE HIV BR
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KW AIDS;
XX Synthe
PN WO9416
XX 26-JAN
PR 11-MAR
PR 11-MAR
PR 11-MAR
PR 11-MAR
PR 11-MAR
PR 15-JUL
PR 26-JAN
PR 11-MAR
PR 11-M
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Best Local
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A genetic vaccine against HIV contains a DNA construct which comprises the sequence encoding gp160. The genetic material was then introduced into the cells of an individual by (a) contacting the individual's cells with a polynucleotide function enhancer (bupivacaine) and (b) administering to the cells the nucleic acid molecule free of retroviral particles. Nucleic acid molecules which are delivered to cells may serve as genetic templates for proteins that function as prophylactic and/or therapeutic immunising agents, replacement copies of defective, missing or non-functional genes; genetic templates for antisense molecules or as genetic templates for antisense molecules or as genetic templates for routeins; genetic templates for antisense molecules or as genetic templates for routeins;
                                                                                                                                                                                                                                                                                                                                                  Method for introducing genetic material into polynucleotide function enhancer and nucleic particles, e.g. HIV immunisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Principal Neutralising Determinant R42161. (Updated on 25-MAR-2003 to 2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (WILL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-JAN-1993;
11-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Weiner DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUL-1993;
21-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-JAN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-AUG-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9416737-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunisation; vaccine; therapy; prophylaxis; defective gene;
non-functional gene; template; antisense; ribozyme; bupivaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV BRU V3 loop peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
21-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR57470;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR57470 standard; protein; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ZURA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MERV/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (WEIN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
les 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunodeficiency virus; acquired immune deficiency syndrome; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CONEY L R.
MERVA M J.
ZURAWSKI V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WEINER D
WILLIAMS
WANG B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RAFVTIGK 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RAFVTICK 8
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                                                                                                                                                                                                                                                                                                      Page 44; 136pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Williams WV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ş
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised). (first ent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93US-00008342.
93US-00029336.
93US-00093235.
93US-00124962.
93US-00125012.
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Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'n
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       motif GPGR are given in AAR42153-
correct PN field.) (Updated on 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Coney LR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB
3.2;
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acid free of reta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Merva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bupivacaine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zurawski VR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                          Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches, 8
                                                                                                          This is the sequence of the V3 loop of the gp120 envelop protein from HIV -1 strain IIIb. The sequence was used with a construct comprising part of the HIV-1 gp120 V3 loop common consensus PND sequence (AAM07390) fused to part of the HIV-1 CD4 binding sitte (AAM07391) and with the V3 loop sequences from HIV-1 strains Thai B (AAM07393) or HGP-30 (AAM07394) to generate a lysine branched peptide which is useful for the prevention and treatment of AIDS. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                       Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                             HIV-1; gp120;
binding site;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  epitope targetted by HIV neutralising antibodies) and was used to determine whether the anti-gp160 antibodies elicited in mice immunised with the genetic vaccine were reactive with this region. (Updated on 29 MAR-2003 to correct PN field.)
                                                                                                                                                                                                    Example 2; Page 5-6; 8pp; Japanese
                                                                                                                                                                                                                                     Novel AIDS vaccine -
                                                                                                                                                                                                                                                             WPI; 1996-461278/46
                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV-1 strain IIIB gp120 V3 loop sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-OCT-2003
24-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW07392;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW07392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 22 AA;
                                                                                                                                                                                                                                                                                                                   27-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                               10-SEP-1996.
                                                                                                                                                                                                                                                                                                                                          27-FEB-1995;
                                                                                                                                                                                                                                                                                  (TERU ) TERUMO C
(OKUD/) OKUDA K.
14
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                                          Similarity
8; Conserv
                                                                                                                                                                                                                                                                                              TERUMO CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                    RAFVTIGK
                                                                                                                                                                                                                          env protein.
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RAFVTICK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
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                                         100.0%;
ilarity 100.0%;
Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                              V3 loop; common consensus PND domain; envelop; CD4; stem-loop; lysine branched peptide; AIDS.
                                                                                                                                                                                                                                                                                                                   95JP-00038835
                                                                                                                                                                                                                                                                                                                                          95JP-00038835
21
                     8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100
                                                                                                                                                                                                                                     comprises branched lysine peptide fragments derived
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                                                     Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 39;
Pred. No.
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                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>,</u>
                                                               Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-
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RESULT 113
AAY07488
ID AAY074
XX

AAY07488

standard;

peptide;

22

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RESULT 114
AAY85137
ID AAY851
XX AAY851
XC AAY851
XX T12-SEP
DT 12-SEP
DT 20-JUN
XX
DE HIV-1
XX
W Human
KW 91ycop
KW V3 loo
XX
PN US6039
XX
PD 21-MAR
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                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents the V3 loop from the gp120 protein of the human immunodeficiency virus-1 (HIV-1) strain IIIB. The invention relates to the generation of heterohybridomas producing human monoclonal antibodies (see AX79204-X79207) to a neutralising epitope of HIV-1 prepared by transforming peripheral blood lymphocytes with Epstein-Barr virus. The antibodies can be used to treat someone infected with HIV-1 or suffering from AIDS. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUN-1990;
12-APR-1991;
23-APR-1992;
                                                                            glycoprotein
V3 loop.
                                                                                    Human immunodeficiency virus type glycoprotein 140; gp140; neutralis
                                                                                                                          HIV-1 IIIB V3 loop peptide sequence.
                                                                                                                                                      12-SEP-2003
20-JUN-2000
                                                 Human immunodeficiency virus
                                                                                                                                                                                              AAY85137;
                                                                                                                                                                                                                    AAY85137 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5; Col 24; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heterohybridoma producing human monoclonal antibodies to immunodeficiency virus-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-370481/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gorny MK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYNY ) UNIV NEW YORK STATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5914109-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              epitope; V3 loop; heterohybridoma; human immunodeficiency v
peripheral blood lymphocyte; Epstein-Barr virus; EBV; AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Light chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV-1 strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-OCT-2003
17-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY07488;
                                                                                                                                                                                                                                                                                                                                      Local Similarity les 8; Conserv
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                                                                                                                                                                                                                                                                                 RAFVTIGK 20
                                                                                                                                                                                                                                                                                                            RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                                                      22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zolla-Pazner S;
                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            variable region; human; HIV-1; gp120; monoclonal antibody; loop; heterohybridoma; human immunodeficiency virus-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IIIB gp120 V3 loop sequence
                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90US-00538451.
91US-00684090.
92US-00872675.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94US-00345321
                                                                                                                                                                                                                                                                                                                                  100.0%; Sc
100.0%; P;
ative 0;
                                                                                      neutralising
                                                                                                                                                                                                                    22
                                                                                                                                                                                                                                                                                                                                                Score 39; I
                                                                                                                                                                                                                                                                                                                                      Mismatches

    HIV-1; infection; prevent; detect;
    antibody; conformational epitope;

                                                                                                                                                                                                                                                                                                                                                              BB
                                                                                                                                                                                                                                                                                                                                                2;
                                                                                                                                                                                                                                                                                                                                                             2;
                                                                                                                                                                                                                                                                                                                                    <u>,</u>
                                                                                                                                                                                                                                                                                                                                                             Length 22;
                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human
                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                   Gaps
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21-MAR-2000. US6039957-A.

21-DEC-1998;

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RESULT 115
ABU07537
ID ABU0753
XX ABU075:
XX ABU075:
XX ABU075:
XX ABU075:
XX CABU075:
XX 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprises administering to a human, a recombinant uncleaved gp140 protein retaining its oligomeric structure. The human produces neutralising antibodies against conformational epicopes of the HIV-1 gp140 protein found on the oligomeric structure of the gp140. The anti-HIV-1 gp140 antibodies of the invention can be used for preventing and diagnosing an HIV infection in a mammal. Gp140 antibodies are useful for treating an HIV infection. A diagnostic method using the antibodies involves isolating a body fluid, preferably blood, and contacting it with a labelled monoclonal antibody for gp140, and detecting any bound antibody. (Updated on 12-SEP-2003 to standardise OS field)
                                                        21-DEC-1998;
                                                                                                                                                                                                                                                                                  Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                             GalNac T4; N-acetylgalactosaminyl transferase T4; acceptor substrate; glycosylation; mucin 1; MUC1; vaccine; antiinflammatory; GalNAc-T1; GalNAc-T2; GalNAc-T3; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents a human immunodeficiency virus type-1 IIIB V3-loop peptide sequence. The peptide sequence is used to test the reactivity of the antibodies of the invention. The invention relates to method for the production of neutralising antibodies against conformational epitopes of HIV-1 envelope proteins in humans. The method
                                                                                                                15-OCT-2002.
                                                                                                                                                                        US6465220-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human N-acetylgalactosaminyl transferase T4,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-OCT-2003
13-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABU07537;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABU07537 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Producing neutralizing antibodies useful for preventing, treating and diagnosing an HIV infection in a mammal comprises administering recombinant uncleaved gp140 proteins to a human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 10; Col 12; 15pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-270121/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Doms RW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                     immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RAFVTICK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first en
                                                           98US-00217306.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97US-00805889
                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                          note= "Gln is acetylated"
                                                                                                                                                                                                                                                           label= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Earl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Broder CC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GalNac T4, substrate #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 116
AAR04502
ID AAR045
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to glycosylating a MUC1 (mucin 1) acceptor substrate, comprising glycosylating the substrate with enzymatically active N-acetylgalactosaminyltransferase (GalNAc)-T1, GalNAc-T2 or GalNAc -T3, or with GalNAc capable of glycosylating MUC1 glycosylation sites that can be glycosylated by GalNAc-T1, GalNAc-T2 or GalNAc-T3, and glycosylating the substrate with enzymatically active human GalNAc-T4 to glycosylate specific Ser, Thr positions in the MUC1 substrate. The method is used for glycosylating an MUC1 acceptor substrate. The glycosylated substrates are useful in preparation of vaccines and antiinflammatory agents. GalNAc-T4 exhibits a different substrate specificity than previously characterised GalNAc transferases. The activity of GalNAc-T4 is unique and specific to glycosylate specific serine and threonine residues in MUC1 tandem repeat. The present sequence is an acceptor substrate peptide used to test the substrate specificity the human GalNac T4 protein, HIVHIBSP120. (Updated on 23-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 8
                                                                                             03-OCT-1988;
01-JUN-1989;
19-SEP-1989;
                                                                                                                                                                                                                                                                                                                               09-SEP-2004
25-MAR-2003
20-SEP-1990
                                                                                                                                                                                                                                                                                                     Cpd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Glycosylating MUC1 acceptor substrate, by glycosylating substrate with acetylgalactosaminyltransferase T1, GalNAc-T2 or GalNAc-T3, then with human GalNAc-T4 to glycosylate specific Ser, Thr residues in substrate.
                         Lynn
                                                                                                                                                                                                                                                            HIV; therapy; prophylaxis.
 WPI; 1990-147824/19
                                         Rusche
                                                                                                                                                   03-OCT-1988;
                                                                                                                                                                              19-APR-1990.
                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                       AAR04502;
                                                                                                                                                                                                                                                                                                                                                                                                               AAR04502 standard; protein; 23 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6;
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                                                                                                                                                                                                        WO9003984-A
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nes 8; Conserv
                                                                                                                                                                                                                                                                                                    eliciting,
                         Ďű,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10
                                      JR,
                                                                 REPLIGEN CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RAFVTIGK 17
                         Putney SD,
Petrobre J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Col 9; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Clausen H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; ilarity 100.0%; Conservative 0;
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                                                                                                                                                                                                                                                                                                                               (revised)
(revised)
(first entry)
                                                                                                                                                                                                                                                                          AIDS; principal neutralising domain;
                                                                                             88US-00252949.
89US-00359543.
89US-00407663.
                                                                                                                                                   88US-00252949
                                                                                                                                                                                                                                                                                                    binding with neutralising antibodies to HIV variants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bennett EP,
                                         Javaherian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 39; DB
Pred. No. 0.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                        ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eisenkraetzer D,
                                        Farley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6
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                                         Grimaila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                           antibodies; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            substrate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 117
AARO4476
ID AARO4476
AC AARO44
XX AARO44
XX O9-SEP
DT 25-WAR
DT 25-WAR
DT 25-WAR
DT WO9-SEP
XX HUMAN
XX HUV is
KW HIV is
KW Princi
XX Synthe
XX WO9003
XX I9-APR
XX 19-APR
XX 19-APR
XX I9-APR
XX I9-APR
XX I9-APR
XX IP-APR
XX I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                       03-OCT-1988;
01-JUN-1989;
19-SEP-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-SEP-2004
25-MAR-2003
20-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bither the N-terminal (a) or C-terminal (b), but not both, may be omitted; either (a) or (b) may comprise any of the following: cysteine, a protein or other moiety capable of enhancing immunogenecity, a peptide from an HIV principal neutralising domain, peptide capable of stimulating T-cells, or general immune stimulant. See also AAR04427-R04506 and AAQ04273-Q04279. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                           Principal neutralising domain of HIV variants - used for producing peptide(8) and antibodies for diagnosis, prophylaxis and/or therapy therapy of HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-OCT-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV isolates HIV-IIIB and HIV-RF; hybrid peptide PR140; therapy; A principal neutralising domain; antibodies; diagnosis; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR04476;
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                                                                                                                                                                                                                                                                    WPI; 1990-147824/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus hybrid peptide RP140
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                                                                                                                                                                                                                                                                                                                                            JR,
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8; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RAFVTIGK 8
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                                                                                                                                                                                                                                                                                                                       Petrobre
                                                                                                                                                                                                                                                                                                                     Putney SD, etrobre J;
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(first entry)
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89US-00359543.
89US-00407663.
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Pred. No.
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Peptide RP140 comprises segments of the Principal Neutralising Domain (envelope protein) from isolates HIV-RF and HIV-IIB. The last Cys residue is added for the purpose of crosslinking to carrier proteins. Cysteine residues may be added, so that the residues at or near both ends

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RESULT 118
AAB66704
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is useful for stimulating an anti-HIV-1 env immune response in a mammal, by stimulating the formation of neutralizing antibodies against conformational epitopes of HIV-1 env protein in a mammal. gp140 is also useful for preventing HIV infection in a mammal. (Updated on 11-SEP-2003 to standardise OS field)
                                                                                                                      The present invention relates to an immunogenic composition comprising a recombinant uncleaved gp140 protein which is a C-terminally truncated form of HIV-1 gp160 protein, missing the gp41 transmembrane domain, and retaining its oligomeric structure, such that neutralizing antibodies against conformational epitopes of HIV-1 envelope proteins found on the oligomeric structure of are produced in an immunized human. The invention
                                                                                                                                                                                                                                                                                                                                                                                Immunogenic composition for stimulating mammalian immune response, comprises recombinant uncleaved gp140 protein retaining its oligomeric structure such that antibodies against HIV envelope proteins are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Earl PL,
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03-MAR-1997;
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                                                                                                                                                                                                                                                                                                               Example 10;
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09-APR-2001
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8; Conserv
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97US-00805889.
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Pred. No.
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Sequence 23

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Human immunodeficiency virus 1

HIV; AIDS; ARC; gp120;

RP135.

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AAR06211
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Best Local
                               Residues 301-324 of HIV gp 120 protein used in polysaccharide by affinity chromatography.
                                                         24-OCT-2003
18-JAN-1991
                                                                                                                                                                                                                                     Protease inhibitor may be use to suppress organ transplant rejection without serious side effects. It may also be used in prevention and therapy of allergy, aplastic anaemia and systemic erythaematodes. Se
                                                                                   AAR07018;
                                                                                                    AAR07018 standard;
                                                                                                                                                                                                                Sequence 24 AA;
                                                                                                                                                                                                                                 also AAR06212
                                                                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                          Protease inhibiting peptide immuno-suppressant - rejection reaction in organs transplantation.
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                                                                                                                                                                                                                                                                          Page 181; 6pp; Japanese.
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Pred. No. 0.21;
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Pred. No.
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25-MAR-2003
28-JAN-1993
Synthetic peptide(s) which enhance infectivity of HIV-1 in cellular cultures - are used for determining HIV-1 virus in blood and other biological materials.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-HIV sulphated polysaccharide (SPS) can prevent syncytium formation in HIV infected C4 cells. SPS may be isolated by affinity chromatography with the given resin bound peptide fragment RP135. (Updated on 24-OCT-2003 to standardise OS field)
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05-JUL-1989;
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                                                                                                                                           De Rossi A, P
Chieco-Bianchi
                                                                                                                                                                                                                                                                                 30-JAN-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . of anti-HIV sulphated polysaccharide - by affinity chromatography a resin-bound peptide corresp. to a HIV gp. 120 fragment.
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CONSIGLIO NAZ DELLE RICER
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8; Conserv
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(revised)
(first entry)
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89US-00375795.
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100.0%; Pr
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Pred. No. 0.21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     strain HTLV-III B
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RESULT 122
AAR29233
ID AAR292
XX AAR292
XX AAR292
XX 25-MAR
DT 14-APR
XX 100
KW V3 100
KW conjug
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              v3 loop; gp41; envelope protein; MN; prototype; virus; varient; HIV; homology; heteroconjugate; enzyme; epitope mapping; replication; conjugate; immunogenic carrier; keyhole limpet hemocyanin; KLH; ovalbumin; succinyl maleimidomethyl cyclohexanylcarboxylate; SMCC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                         Hetero-conjugate antibodies for treating HIV infections antibody specific for an effector cell surface antigen and
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14-APR-1993
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                                                                                                                                                                                                                                                                                                                                                                           Higgins PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-MAY-1991;
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                                                                                                                                                           Disclosure; Page 19; 69pp; English
                                                                                                                                                                                                                                                                                                                            WPI; 1992-415475/50.
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                                                                                                                                                                                                                of GP-120 envelope
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(first entry)
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Pred. No.
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The sequences given in AAR29226-35 represent peptides which were used as immunogens for the production of antibodies against HIV. These peptides may be either unconjugated or conjugated to an immunogenic carrier, eg. keyhole limpet hemocyanin (KLH) or ovalbumin, using succinyl maleimidomethyl cyclohexanylcarboxylate (SMCC) as a conjugating agent.

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RESULT 123
AAR26870
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Best Local :
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                                                                                                     The sequence is that of peptide IIIB, derived from the V3 region of HIV gp120, it was used in binding assays for anti-gp120 antibodies. The anti-gp120 antibodies are useful in vaccine formulations for the treatment or prevention of HIV infection. See also AAR26867-R26873. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          heteroconjugate enzymes of the invention. The antibodies raised by the these sequences may be identified by standard epitope mapping techniques. These antibodies are capable, even at low concentrations, of nearly eliminating viral replication of different strains of HIV. (Updated on 25 -MAR-2003 to correct PN field.)
                                                                                      Sequence
                                                                                                                                                                        Example; Page 9; 30pp; Japanese.
                                                                                                                                                                                            Anti-idiotype antibodies and methods for their selection - useful vaccines for the prevention and treatment of HIV infection.
                                                                                                                                                                                                                              WPI; 1992-309988/38
                                                                                                                                                                                                                                                                          (IDEC-) IDEC
                                                                                                                                                                                                                                                                                             11-MAR-1991;
06-MAR-1992;
                                                                                                                                                                                                                                                                                                                               11-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                   16-SEP-1992.
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                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus; AIDS; anti-gp120 antibodies.
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20-MAY-1998
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                    RAFVTIGK 8
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                                           Conservative
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92US-00894766.
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Pred. No.
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Pred. No. 0.21;
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                                         Mismatches
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AAR32406 ID AAR

AAR32406 standard; peptide;

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RESULT 124

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RESULT 125
AAR33190
ID AAR331
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Matches 8
                          Synthetic
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                                                                                                 25-MAR-2003
11-JUL-1993
                                                                                                                                                                                                                                                                                                                                              Nine peptides from the V3 regions of HIV-1 isolates IIIb, RF and MN were incorporated into tetravalent multiple antigen peptide systems (MAPS) (see ARA32406-14). Parallel groups of three peptides with chain lengths spanning from 11-24 residues were synthesised in MAPS formate for each isolate. ELIS assays demonstrated that antisera titers in mice were closely related to the length of the IIIb peptide used for the immunisation - the longer the stronger the response. There was no substantial antibody prodn. in mice against the other two series of peptides, RF (B4-B6), and MN (B7-B9), except for a low reactivity in the gp. immunised with B8 (MN isolate). Specificity tests of the B cell response demonstrated that the T cell epitope (AAR32415) also serves as B cell epitope. (Updated on 25-MAR-2003 to correct PN field.)
                                                 AIDS; HIV;
                                                                                                                                       AAR33190;
                                                                                                                                                             AAR33190 standard; peptide;
                                                                                                                                                                                                                                                                                                                           Sequence 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Fig 1; 35pp; English.
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                                              therapy; autoimmune disease; gp120;
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Pred. No. 0.21;
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equence IGPGR.
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RESULT 126
AAR38165
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Best Local S
Matches 8
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25-MAR-2003
12-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Two peptides, gp41[8] and V3 loop derived from the gp120 region of HIV-1 were synthetised and purified. To improve the solubility of the gp41[8] peptide the sequence RSS was added to the amino terminal to produce peptide R-S-Sgp41[8]. The immunodominant HIV-derived peptides were used to ascertain whether E.coli outer membrance protein TraT augments the in vitro T-cell proliferative responses. (Updated on 25-MAR-2003 to correct PN field.)
Peptide composition for treating and preventing viral infections comprise CTL-inducing epitope and HIV infection-inhibiting sequentely the composition of the composi
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                                                                                                                                                                                                                                                                    02-DEC-1991;
16-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gp120; HIV-1; cytotoxic T-lymphocyte; CTL; T-helper; AIDS; infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             V3 loop peptide N24G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Compsns. contg. E.coli outer membrane proteins TraT, OmpA or OmpF - increase immune response and are used for treating auto:immune diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-AUG-1992;
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                                                                                                                                                            Sastry JK,
                                                                                                                                                                                                                                                                                                                                                 02-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9310816-A1
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                                                                                                                                                                                                                                                                                                                                                                                                     10-JUN-1993.
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                                                                                                                                                                                                                   (TEXA ) UNIV TEXAS SYSTEM
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8; Conserv
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                                                                                                                                                            Arlinghaus RB,
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(revised)
(first entry)
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92US-00945865.
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                                                                                                                                                                 Platsoucas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bell SJD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cooper DA;
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RESULT 127
AAR44191
ID AAR441
XX AAR441
AC AAR441
XX 25-MAR
DT 25-MAR
DT 20-MAX
XX Upp120
XX Antige
KW Antige
KW Strain
KW Lipoph
KW Cytcoto
KW Pathogo
XX Synthe
XX Synthe
XX WO9322
XX Human
OS Synthe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV gp120 V3 loop-derived peptides (AAR38170-87) are successful in generating CTL responses, esp. peptide R15K (AAR38167); the T-helper cell inducing peptide includes the sequence C19A (AAR38164); HIV infection-inhibiting peptides are given in AAR38188-206, and are esp. peptides R15K, N24G, E13V, R8K, T13Q and H13N (AAR38165-69). The peptides may also be derived from an influenza virus protein or a sendai virus protein (AAR41014-15). It was observed that peptide N24G (amino acids 308-311), with sequences derived from the V3 loop of HIV-1 IIIB, inhibits HIV-1 infection of primary human T cells by 92% at 1 microg/ml (ca. 0.4-0.6 microm). (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG
                                                                                                         New multiple antigen system esp. for lipophilic membrane anchor imparting antigens coupled to dendritic core.
                                                                                                                                                                                                                        WPI; 1993-368723/46.
                                                                                                                                                                                                                                                                                                                                                                                        01-MAY-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                        03-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antigen; B2; third variable domain; V3 loop; gp120; HIV-1; vaccine; strain IIIB; multiple antigenic peptide system; dendritic core; lipophilic membrane anchoring group; mammal; humoral; immunisation; cytotoxic T_cell; CT; immune response; infection; Freunds adjuvant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gp120 V3
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20-MAY-1994
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                                                      Disclosure; Fig 8; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9322343-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pathogen; HIV; influenza; malaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR44191 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide
                                                                                                                                                                                                                                                                                                                                  (UYRQ )
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                                                                                                                                                                                                                                                                                                                                       VIND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             loop antigen B2 and lipophilic membrane anchoring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RAFVTIGK 8
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              93WO-US004179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Lipophilic membrane anchoring group"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              label= B2 antigenic peptide
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Pred. No.
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                                                                                                                                   use in HIV vaccines adjuvant activity, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2
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                                                                                                                                   contains
d peptide
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The sequence given in AAR44190 is a peptide antigen, B2,

which represents

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RESULT 128
AAR63821
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Best Local
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AAR63809-R63849 are epitopes from the human immunodificiency virus type 1 (HIV-1) gp120, by binding one or more of these epitopes to a carrier a HIV vaccine is produced. These vaccines can elicit the production of HIV-neutralising antibodies in monkeys, and therefore may be used to prevent HIV infections, and to highten the immune response in HIV infected humans. (Updated on 25-MAR-2003 to correct pN field.) (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                           Claim 1; Page 18; 77pp; English
                                                                                                                                                                                                                                             New peptide(s) comprisi against HIV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             residues 312-329 of the third variable domain (V3 loop) of gp120, of HIV-
1 strain IIIB. This sequence was attached to an amino acid linker (see
also AAR44191) in the production of a multiple antigenic peptide system.
This system comprises a dendritic core to which are covalently attached
at least one peptide, eg. an antigenic peptide, and a lipophilic membrane
anchoring group. This system may be injected into a mammal and elicits
both humoral and cytotoxic T cell (CTL) immune responses. This system may
be used to immunise against HIV infection. The lipophilic membrane
anchoring group provides efficient adjuvant activity without the toxicity
problems of Freunds adjuvant, while the dendritic structure allows
                                                                                                                                                                                                                                                                                                                     WPI; 1994-341488/42
                                                                                                                                                                                                                                                                                                                                                                  Vahlne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-OCT-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV neutralising antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1; HIV-1; gp120 epitopes; vaccines;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV-1 gp120-24 epitope amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-OCT-2003
25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          multiple antigens to be attached. Optionally the antigens may be derived from different pathogens, providing vaccines which protect against more than one disease eq. HIV, influenza and malaria. (Updated on 25-MAR-2003)
                                                                                                                                                                                                                                                                                                                                                                                                               (SYNT-) SYNTELLO VACCINE DEV AB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-JUN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                               Svennerholm B,
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                                                                                                                                                                                                                                                                 comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                              93US-00048976.
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                                                                                                                                                                                                                                                                 HIV gp120
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Pred. No. 0.21;
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                                                                                                                                                                                                                                                                 epitope(s) -
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Query Match

100.0%;

Score

39;

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Length

24

25-JAN-1999

(first entry)

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Sequence

24

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AAW67414
ID AAW6
XX
AC AAW6
XX
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AAR74608
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                                                                                                                                                             Query Match
Best Local
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                                                                                                                                                                                                                      The peptide represented in this sequence, and those represented by sequences AAR74604-7 are epitopes of HIV-1 gpu20 that are effective to generate antibodies that inhibit infection of mucosal cells by HIV-1. These peptides are administered to the epithelium in a vaccine, or are used to generate mucosal antibodies and thereby inhibit infection by HIV-1. These peptides are used for inhibiting the entry of HIV into vaginal and rectal mucosal epithelium. The antibodies that can be generated from them are able to block subsequent infection by HIV. (Updated on 16-OCT-
                                 AAW67414;
                                                  AAW67414 standard; peptide;
                                                                                                                                                                                             Sequence 24
                                                                                                                                                                                                                                                                                                                          Claim 2;
                                                                                                                                                                                                                                                                                                                                                                    HIV-1
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-178653/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV-1; HIV; AIDS; gp120; mucosal cell; epithelium; vagina; rectum; antibody; mucosal administration; vaccine; infection.
                                                                                                                                                                                                                                                                                                                                                                                                             Czerkinsky C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV-1 gp120 peptide #5.
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04-JAN-1996
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                                                                                                                                                                                                                                                                                                                                                          peptide
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                                                                                                                                                 100.0%; ilarity 100.0%; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                        23; 34pp;
                                                                                                                                                                                                                                                                                                                                                                                                           Holmgren
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                                                                                                                                                                                                                OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                      inhibit mucosal epithelium cell in inhibit HIV-1 infection of vaginal
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                                                                                                                                                            Score 39; DB 2;
Pred. No. 0.21;
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                                                                                                                                                                     Length 24;
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al or rectal
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RESULT 131
AAW98904
ID AAW9890
XX AAW989
XX AAW989
XX HIV-1
XX HIV-1
XX Gag pr
XX Synthe
OS Human
XX Synthe
OS Human
XX US5876
XX OS-JUN
XX OS-JUN
PR 09-JUN
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-cell epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitopes are generally designed based on the p24 core protein and the B-cell epitopes from the V3 loop of the gp120 protein from various HIV-1 epitopes from the V3 loop of the gp120 protein from various HIV-1 epitopes.
                                                09-JUN-1993;
09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 24 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic human immunodeficiency virus-1 peptide(s) - containing T-cell epitope and B-cell epitope(s) are candidate vaccines against HIV-1.
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09-JUN-1994;
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Human immunodeficiency virus
                                                                                                                             05-JUN-1995;
                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                         gag protein;
                                                                                                                                                                                                                                                                                                                                                                                      HIV-1; human
                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV-1 vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW98904;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW98904 standard; peptide; 24 AA
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                                                                                                                                                                                   02-MAR-1999
                                                                                                                                                                                                                                US5876731-A.
                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
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  (CONN-) CONNAUGHT LAB LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                         immunodeficiency virus; vaccine; T-cell epitope; B-cell epitope; gp41 protein; chimeric; infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                    synthetic peptide SEQ ID NO:99.
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94US-00257528.
                                                93US-00073378.
94US-00257528.
                                                                                                                             95US-00462507.
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Pred. No. 0.21;
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RESULT 132
AAY22581
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV LDL binding peptide, sequence A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-OCT-2003
19-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chong
                                                                                                                                                                                                                                    Enhancing the immune response using a recombinant human low-density lipoprotein receptor, useful for treating viral infections, especially human immunodeficiency virus (HIV) infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV-infected CD4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention
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This sequence represents a HIV sequence that binds human low density lipoprotein (LDL), and is designated sequence "A". The invention relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-JAN-1999;
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                                                                                                                                       Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-494040/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PREN/) PRENDERGAST P T.
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(first entry)
                                                                                                                                       19; 24pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-0072980P
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₿
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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RESULT 133
AAY22583
ID AAY22583
XX AAY225
XX AAY225
XX IT-OCT
DT 19-OCT
DT 19-OCT
XX ITMUNO
KW HIV: L
KW ITWUNO
KW VITAL
KW ACQUIA
KW AC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC to a method for enhancing the immune response in a patient with a CC condition, selected from immunodeficiency (due to a viral, bacterial, CC mycoplasmic, fungal or parasitic infection, or from the growth of CC neoplastic tissue), myalgic encephalomyelitis (ME), post inoculation or CC viral infection fatigue syndrome, tuberculosis, or hepatitis. The method CC comprises using a pharmaceutical composition, comprising a recombinant CC thuman LDL receptor or a mimic molecule to the cysteine rich domain of LDL receptor. The human recombinant LDL receptor forms pharmaceutical CC compositions for: the treatment of acquired immunodeficiency syndrome (AIDS) or ARC (AIDS related complex); reducing syncytium formation in HIV -infected CD4 cells; treating blood or body fluid or organs to neutralise/remove immunosuppressive peptides and/or viruses; or treating CC neutralise/remove immunosuppressive peptides and/or viruses; or treating conditions a human or animal host. The human recombinant LDL receptor is also useful for manufacturing medicaments for treating all the conditions given above. The human recombinant LDL receptor is a highly specific inhibitor of HIV-1 replication in vitro. (Updated on 17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Loc
Matches
                                                                                                                                                                                                                                   Enhancing the immune response using a recombinant lipoprotein receptor, useful for treating viral in human immunodeficiency virus (HIV) infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV; LDL; low density lipoprotein; human; immune response; infection; immunodeficiency; neoplastic tissue; myalgic encephalomyelitis; ME; viral infection fatigue syndrome; tuberculosis; hepatitis; AIDS; ARC; acquired immunodeficiency syndrome; AIDS related complex; HIV-infected CD4 cell; immunosuppressive peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prendergast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-AUG-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-494040/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV LDL binding peptide, sequence "A" variant.
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19-0CT-1999
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8; Conserv
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Pred. No. 0.21;
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                                                                                                                                                                                                                                                                                                            human low-density
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This sequence represents a variant of the HIV sequence that bind low density lipoprotein (LDL), and is designated sequence "A" (se AAY22581). The sequence "A" peptide is isolated from HIV isolate IIIB (BH10), and this sequence was isolated from HIV isolate IIIB

at binds human "A" (see

This sequence represents a fragment of a HIV1 protein, and can be use the immunogenic composition of the invention. The composition compris synthetic fusion polypeptide which includes a sequence encoding 1 or T cell epitopes and a sequence encoding 1 or more B cell epitopes and carrier. Both the T cell and B cell epitopes are derived from HIV

and can be used in

from HIV

more d a

infection

6

73-74;

43pp; English.

IIIB(BH8)

Disclosure; Page 12;

24pp; English

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RESULT 134
AAY39769
ID AAY397
XX AAY397
XX AAY397
XX 17-OCT
DT 26-NOV
XX HIV; v
KW HIV; v
KW infect
XX US5951
XX INFECT OCCUN-
PH O6-JUN
PR 09-JUN
PR 10-JUN
XX CONN-
PT CONTAI
PT VITUS,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC The invention relates to a method for enhancing the immune response in a CC patient with a condition, selected from immunodeficiency (due to a viral, CC bacterial, mycoplasmic, fungal or parasitic infection, or from the growth CC of neoplastic tissue), myalgic encephalomyelitis (ME), post inoculation CC or viral infection fatigue syndrome, tuberculosis, or hepatitis. The CC method comprises using a pharmaceutical composition, comprising a CC recombinant human LDL receptor or a mimic molecule to the cysteine rich CC domain of LDL receptor. The human recombinant LDL receptor forms CC pharmaceutical compositions for: the treatment of acquired compositions cc gyncytium formation in HIV-infected CD4 cells; treating blood or body cluid or organs to neutralise/remove immunosuppressive peptides and/or CC fluid or organs to neutralise/remove immunosuppressive peptides and/or CC recombinant LDL receptor is also useful for manufacturing medicaments for treating all the conditions given above. The human recombinant LDL receptor is a highly specific inhibitor of HIV-1 replication in vitro. CC (Updated on 17-OCT-2003 to standardise OS field)
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                             Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency virus, useful antigens in producing vaccines.
                                                                                                                                                                                                                                           WPI; 1999-550482/46.
                                                                                                                                                                                                                                                                                                                                                     09-JUN-1993;
09-JUN-1994;
                                                                                                                                                                                                                                                                             Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                         06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5951986-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV; vaccine; immunogenic composition; T infection; antibody; antiviral.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV; vaccine;
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26-NOV-1999
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                                                                                                                                                                                                                                                                                                                (CONN-) CONNAUGHT LAB LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimeric
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8; Conserv
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                                                                                                                                                                                                                                                                         Chong P,
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ilarity 100.0%;
Conservative
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(first en
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94US-00257528.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                             Sia CDY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cell epitope; B cell epitope;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mmune response in a ncy (due to a viral, or from the growth post inoculation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Best Local (
                                                       The present invention concerns the identification of a number of chemokines which can be used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences ABB15785-B15794, ABB15803-B15813 and ABB15831-B15848. These chemokine derivatives can be used to treat diseases such as autoimmune diseases, atherosclerosis, osteoporosis, HIV infection and AIDS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotoxaemia, myocardial infarction, acute ischaemia and rheumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in
                                                                                                                                                                                                                                                               New peptide 3, amide and heterocyclic compounds and saccharide conjugates used for inhibiting chemokine induced activity and for treating e.g. stroke, vascular diseases, autoimmune diseases and tumor growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                monocyte chemoattractant protein-1; inflammation; atherosclerosis; HIV; AIDS; stroke; psoriasis; autoimmune disease; hypertension; endotoxaemia; basophil-mediated disease; myocardial infarction; acute ischaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                opsonising and antiviral. The peptide components may be selected to induce a response against different viral isolates and in subjects who recognise different T cell epitopes. (Updated on 17-OCT-2003 to
                                                                                                                                                                                                                                                                                                                               WPI; 2000-499101/44.
                                                                                                                                                                                                                                                                                                                                                             Grainger DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB15873 standard; peptide; 24
                                             gene therapy
                                                                                                                                                                                                                                  Disclosure; Fig 18; 387pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JAN-2000; 2000WO-US000821.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-JUL-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200042071-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Macrophage recruitment; chemokine derivative; MCP-1; osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human chemokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB15873;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The composition induces HIV-1-specific polyclonal antibodies that
                                                                                                                                                                                                                                                                                                                                                                                              (NEOR-) NEORX CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14 RAFVTICK 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   arthritis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                             for the same diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                              Tatalick LM;
                                                                                                                                                                                                                                                                                                                                                                                                                           99US-00229071.
99US-00271192.
99US-00452406.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               derived peptide #25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               contraception
                                                                                                                                                                                                                                   English.
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                                             as
                                             well
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                                             in the production
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Query Match

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Score

39;

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Length 24;

Sequence 24 AA;

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RESULT 136
ARB68602
ANB6866XX
ARB686
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ARB686
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ARB6866
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                                                                                                                                                                                                                                                                                                                 The present invention relates to a liposome composition comprising CC liposomes having an outer surface layer of polyethylene glycol chains, CC each having a free distal end. A polypeptide or polysaccharide effector CC molecule is covalently attached to a portion of the distal ends. The CC effector interferes with specific binding of pathogen or cell in a CC bloodstream to a target cell or cell matrix, and is rapidly removed by CC renal clearance from the bloodstream when administered in free form. The CC binding a pathogen or cell in the bloodstream, to a target cell or cell condition mediated by binding a pathogen or cell in the bloodstream, to a target cell or cell conficient in treating a pathogen or cell in the bloodstream, to a target cell or cell conficient in the present gequence is a peptide of the V3 loop of HIV envelope protein gp120. This CC peptide may be used in the composition of the present invention. gp120 binds to the CD4 receptor during HIV infection of lymphocytes. By CC introducing the present peptide, the CD4 receptors are blocked, thereby CC inhibiting HIV infection. (Updated on 11-8EP-2003 to standardise CS
                                                                                                                                          Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.0%; I Matches 8; Conservative 0;
                                                                                                                                                                                                                                        Sequence 24 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 13; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liposome composition for use in treating septic shock comprises liposomes having an outer surface layer of polyethylene glycol chains, and a polypeptide or polysaccharide effector molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-MAR-1993;
29-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV gp120 V3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-SEP-2003
25-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB68602 standard; peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-201897/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US6180134-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SEQU-) SEQUUS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gp120 V3
15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 RAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 RAFVTIGK 8
                                                                                                                    Similarity
8; Conserv
RAFVTIGK 22
                                                  RAFVTIGK 8
                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Woodle MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      loop; liposome composition; HIV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 loop peptide #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93US-00035443
94US-00316436
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                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Martin FJ,
                                                                                                                       0
                                                                                                                                                Score 39; DB 4;
Pred. No. 0.21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 0.:
; Mismatches
                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barenholz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.21;
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                                                                                                                                                                             Length
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RESULT 138
AAP90281
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AAP82464
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Best Local S
Matches 8
                                                                                       09-SEP-2004
24-OCT-2003
25-MAR-2003
22-JUN-1990
08-MAR-1989
                               Simian-Human
Unidentified
               EP306219-A.
                                                        HIV; AIDS;
                                                                        Peptide 135
                                                                                                                                 AAP90281;
                                                                                                                                                AAP90281 standard; protein; 25
                                                                                                                                                                                                                                                                        This peptide is a component of an AIDS vaccine. It can fold as a maximally amphipathic helix and is recognised by T-cells immune to AIDS virus envelope protein. See also AAP82462-63 and AAP82465-79. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                           Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                   Claim 9; Page 10; 16pp; English.
                                                                                                                                                                                                                                                                                                                                Synthetic peptide(s) as vaccines for AIDS - selected from peptide regions which can fold as a maximally amphipathic helix recognised by I cells.
                                                                                                                                                                                                                                                                                                                                                            WPI; 1988-184640/27.
                                                                                                                                                                                                                                                                                                                                                                            Delisi C,
                                                                                                                                                                                                                                                                                                                                                                                            (USDC )
                                                                                                                                                                                                                                                                                                                                                                                                                    30-DEC-1986;
12-FEB-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                             23-DEC-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JUL-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
12-NOV-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAP82464 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AIDS vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide component of AIDS
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                                                                                                                                                                                                                         Similarity 100
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                            SD
                                                                                                                                                                                                         RAFVTICK 8
                                                                                                                                                                                         RAFVTIGK 16
                                                        env gene; HIV vaccine; ds.
                                                                                                                                                                                                                                                                                                                                                                           Margalit H,
                                                                                                                                                                                                                                                                                                                                                                                          SEC OF COMMERCE.
DEPT HEALTH & HUMAN SERVICE.
                                                                       of HIV env gene
                                      immunodeficiency virus
                                                                                                        (revised)
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(first en
                                                                                       (first entry)
                                                                                                 (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T-cells.
                                                                                                                                                                                                                                                                                                                                                                                                                   86US-00947935.
87US-00014430.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein;
                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                           Cornette JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vaccine.
                                                                                                                                                                                                                         0;
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                                                                                                                                                                                                                                Score 39; DB 1, Pred. No. 0.22;
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                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                          Ouyang
                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                           cs;
                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                       Length 25;
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RESULT 139
AAR04475
ID AAR044
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Best Local Similarity
Thes 8; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rusche Ji
Lynn DU,
                                                                                                         03-OCT-1988;
01-JUN-1989;
19-SEP-1989;
                                                                                                                                                                                                                                                                       09-SEP-2004
25-MAR-2003
20-SEP-1990
Principal neutralising domain of HIV variants - used for producing peptide(s) and antibodies for diagnosis, prophylaxis and/or therapy therapy of HIV infection.
                                       WPI; 1990-147824/19.
                                                                                                                                                                                                                     HIV isolates HIV-IIIB and HIV-RF; hybrid peptide PR137; therapy; AIDS; principal neutralising domain; antibodies; diagnosis; prophylaxis.
                                                                                      (REPK ) REPLIGEN CORP
                                                                                                                                              03-OCT-1988;
                                                                                                                                                                  19-APR-1990
                                                                                                                                                                                    WO9003984-A
                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                    Human immunodeficiency virus hybrid peptide RP137.
                                                                                                                                                                                                                                                                                                               AAR04475;
                                                                                                                                                                                                                                                                                                                                 AAR04475 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protein derivative stimulates a lymphocyte proliferative response infected humans, providing a means of diagnosis, protection and therapeutic value. (Updated on 25-MAR-2003 to correct PR field.) (on 25-MAR-2003 to correct PA field.) (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Revised record issued on 09-SEP-2004 :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New HIV proteins and I therapy of AIDS, esp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rusche JR, Putney SD, Petro J, Okeeffe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1989-070387/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-AUG-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-AUG-1988;
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                                                                                                                                                                                                                                                                                                                                                                                15 RAFVTIGK 22
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                                                        , Putney |
Petrobre
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                                                                                                                                                                                                                                                                                 (revised)
                                                                                                       88US-00252949.
89US-00359543.
89US-00407663.
                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                              88US-00252949.
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Pred. No.
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of vaccines against
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                                                                 Farley J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Correction to location
                                                                                                                                                                                                                                                                                                                                                                                                                             0.22;
                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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                                                                  Grimaila
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Claim 8

(88);

Page 76;

108pp; English.

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RESULT 140
AARO8276
ID. AARO82
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Best Local S
Matches 8
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                            This peptide is derived from the HIV IIIB isolate and is cross- reactive with the HIV major neutralisation determinant (MNtD). This MNtD is used in a conjugate, covalently linked to the outer membrane protein (Omp) from Neisseria, as an immunogen for vaccination against AIDS. A cocktail
                                                                                                                                   Conjugate immunogen for AIDS and determinant of HIV and Neisseria
                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV peptide fragment
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                                                                                                         Claim
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                                                                                                                                                                                                                                                                                                                    06-JUN-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; conjugate immunogen; Neise neutralisation determinant
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89US-00362177.
89US-00362178.
89US-00362179.
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                                                                                                         24pp; English.
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100.0%; P7
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               peptides
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Pred. No. 0.2
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                                                                                                                                                                                                                                                                        The peptide corresponds to residues 294-318 of the gp120 envelope protein of HIV-1 which is a principal neutralising determinant (PND). Abs recognise residues 294-308 (MAb BAT267) or 304-318 (MAb 123). These MAb are used to raise anti-idiotypic Abs (AAbs). The Abs are useful for passive immunisation and as components for immunotoxins which destroy T-cells infected with HIV. They inhibit T-cell infection and syncytium formation, are group specific and neutralise specific strains of HIV-1. They can be used to treat AIDS or ARC. The AAbs can be used for active immunisation or can be admin with another vaccine to increase antigenicity. See also AAR13121. (Updated on 24-OCT-2003 to standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-idiotypic; antibody; gp120; HIV; human immunodeficiency virus; paratope; complemetarity determining region; CDR; immunisation; vacimmunotoxin; T-cell; AIDS; ARC.
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OS field)
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12-JUN-1990;
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RAFVTIGK 25
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                                                                                                Conservative
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90US-00531789.
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RESULT 142

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XX HIV-
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RESULT 143
AAR31276
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Best Local (
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EP467700-A
                                                                                                                                                                                                                                                                         HIV principal determinant peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This peptide is one example of an HIV-1 amplifier peptide for use in a composition for enhancing the immunogenicity of an envelope glycoprotein of a virus. The sequence corresponds to the major neutralisation epitope (loop V3) of HIV-1 bruIII isolate and enhances the induction of persistent neutralising antibodies in the host. The amplifier peptide is used in addition to an envelope glycoprotein for priming the induction of neutralising antibodies. The compositions are particularly useful for vaccinating against HIV, SIV, HTLV-I and HTLV-II
                                                                                                                                                     Synthetic
                                                                                                                                                                                        AIDS; ARC; human immunodeficiency virus; vaccine; Neisseria; meningitidis b; outer membrane protein complex; OMPC; PND135
                                                                                                                                                                                                                                                                                                                                  12-FEB-1993
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                                                                         Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enhancing immunogenicity of envelope glyco:protein - for use as vaccine or immuno:therapeutic drug especially against HIV, HTLV-I and HTLV-II.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
les 8; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 RAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RAFVTICK 8
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                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                              Location/Qualifiers
                                              /note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 39;
100.0%; Pred. No.
                                "bonds to the OMPC of the conjugate via this site"
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The invention relates to a co-conjugate comprising an immunogenic protein or protein complex having a first set of covalent linkages to low molecular weight moieties which have an anionic or polyanionic character at physiological pH, and a second set of covalent linkages to peptides comprising HIV principal neutralizing determinants (PND's) or immunologically equivalent peptides. Preferably at least one set of the covalent linkages is comprised of maleimide derivatives; the (poly)anionic molety is composed of one to five residues of the anionic form of a carboxylic, sulphonic or phosphonic acid; the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria
                                                                  meningitidis b; and the PND peptide has a linear structure, a disulphide-bonded cyclic structure, an amide-bonded cyclic structure or a thioether-bonded cyclic structure. The present sequence (PND135) is an example of a PND peptide component used in the co-conjugate. The co-conjugate is useful for inducing anti-peptide immune response in mammals, for inducing HIV-neutralizing antibodies in mammals, for formulating vaccines to
Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUL-1990;
19-JUL-1990;
19-JUN-1991;
                                    prevent HIV infection or disease, include afflicted with HIV infection or disease
                                                                                                                                                                                                                                                                                                                                                                                                                                    Conjugate proteins comprising HIV peptide components - useful for preparing vaccines for e.g. AIDS or for treating infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-026505/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUL-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUN-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                Page 56; 63pp;
ΑA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Marburg S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90US-00555339.
90US-00555966.
91US-00715276.
91US-00715278.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                English
                                                    including AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Emini
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EA;
                                                formulating vaccines to AIDS, or for treating humans
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AAR30031
ID AAR
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                                                                                                                                                                                 RESULT 144
                                                                                                                                                                                                                                         Best Loc
Matches
                                                                                                                                                                                                                                                  Query Match
Best Local :
                                                                     Human immunodeficiency virus; AIDS; PND; MIEP; conjugate; major immune enhancing protein; vaccine; anti-HIV antibodies; immunogen;
                                                                  passive immunisation.
                                                                                                  HIV principle neutralising determinant 135
                                                                                                                                             AAR30031;
                                                                                                                    25-MAR-2003
28-APR-1993
                                                                                                                                                             AAR30031 standard; peptide; 25 AA
                                                                                                                                                                                                         15
                                                                                                                                                                                                                        1 RAFVTICK 8
                                                                                                                                                                                                                                         Similarity
8; Conserv
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                                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                     100.0%; 5c.
100.0%; Pr
                                                                                                                                                                                                                                                Score 39; DB 2
Pred. No. 0.22;
                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                        2
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                                                                                                                                                                                                                                                       Length 25;
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11-JUN-1992; 23-DEC-1992.

92EP-00201693

Human immunodeficiency virus.

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RESULT 145
AAR26712
ID AAR267
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The peptide is HIV principle neutralising determinant (PND) 135 and is used as part of a conjugate comprising the major immune enhancing protein (MIEP) of Neisseria meningitidis covalently linked to the HIV PND. The conjugate may be used to prepare vaccines against HIV infections, e.g. AIDS, as research tools for studying PND structure- function relationships, or as immunogens for use in the passive immunisation of humans. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 59; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conjugates of HIV-1 PND peptide(s) meningitidis - useful as a vaccine infection, e.g. AIDS in humans.
HIV peptide-polysaccharide-protein conjugates - used in produce antibodies to prevent or treat HIV infection.
                                                                                                                                                                                                                                                        Modified-site
                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                           Human
                                                                                                                                                                                                                                                                                                                                                  HIV-PND-polysaccharide-protein conjugate vaccine
                                                                                                                                                                                                                                                                                                                                                                         09-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                              AAR26712;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 25 AA;
                                    WPI; 1992-034437/05
                                                        Marburg
                                                                                                                19-JUL-1990;
19-JUN-1991;
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                                                                                                                                                                                                          EP468714-A.
                                                                                                                                                                                                                                                                                                                outer
                                                                                                                                                                                                                                                                                                                                                                                                                   AAR26712 standard; peptide;
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                                                                                                       19-JUN-1991;
                                                                                                                                       19-JUL-1990;
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                                                                                (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                immunodeficiency virus; principal neutralizing determinant;
membrane protein complex; OMPC; Neisseria; AIDS; PND135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15
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8; Conserv
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                                                         Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                    90US-00555558.
90US-00555974.
91US-00715275.
91US-00715277.
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                                                                                                                                                                                                                                                       Location/Qualifiers
1. .1
                                                                                                                                                                                                                                            /note= "Joins onto polysaccharide-protein complex via
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                                                         Emini
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Pred. No. 0.22;
                                                         EA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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RESULT 146
AAR33222
ID AAR332
XX AAR3322
XX AAR332
XX IS MAI V 2
XX IS MAI V 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a conjugate of an HIV principal neutralizing determinant (PND), or an immunologically equivalent peptide (PEP), covalently coupled to an immunologenic protein or protein complex through an anionic polysaccharide linker. Pref. the immunogenic protein is the outer membrane protein complex (OMPC) of Neisseria meningitidis b and the PND peptide has a linear structure, a disulphide-bonded cyclic structure, an amide-bonded cyclic structure or a thioether-bonded cyclic structure. The present sequence (PND135) is an example of a PND peptide component. The conjugates are used for inducing HIV-neutralising antibodies or for making vaccines to prevent contraction of HIV infection or disease. The antibodies can be used for passively protecting against infection by HIV. or for protecting against proliferation of HIV post-infection, or for treating AIDS, or in diagnostic assays
                                                                                                                                                                                  Synergistic compsn. f
V3 loop of GP120 and
polypeptide.
                                                                                                                                                                                                                                                                                           WPI; 1993-100653/12.
                                                                                                                                                                                                                                                                                                                                              Potts BJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                          09-SEP-1991;
20-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV gp120 V3 loop immunogenic peptide RP135 (IIIB).
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13-JUL-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9304693-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            infection;
                                                                                                                                                                                                                                                                                                                                                                                               (REPK ) REPLIGEN CORP
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8; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25 AA;
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                                                                                                                                                                                                                                                                                                                                              Whitescharf ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CD4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunodeficiency virus;
D4 binding site; soluble
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                91US-00756677.
92US-00916542.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92WO-US007511.
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25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "not in natural sequence of isolate"
                                                                                                                                                                                                             for treating HIV-1 infection - comprises antibody dantibody to CD4 binding site of GP120 or soluble
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                                                                                                                                                                                                                                                              Field KG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       y virus; antibody generation; AIDS; soluble CD4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                              Herlihy WC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.22;
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The sequence is that of peptide RP135 (IIIB) used as an immunogen for generation of antibodies directed against the V3 loop of HIV gpl20. The antibodies can be used as part of a compsn. With antibodies directed against the CD4 binding site of gpl20. The antibodies act synergistic

act synergistically

tor the

Example; Page 12; 56pp;

English

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AAR41336
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Matches 8
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Matches 8
                                                                                                              The sequences given in AAR41336-39 and AAR42664 represent peptides derived from the V3 region of HIV envelope gp120. These peptides may be used in a vaccine which is effective in humans and animals and activates humoral and cellular immunity. The vaccine also contains a carrier protein containing a cysteine group, eg. human serum albumin (HSA), keyhole limpet haemocyanin (KLH) or multiple antigen peptide. (Updated or 25-MAR-2003 to correct PN field.) (Updated on 24-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                        Virus for prevention of HIV infected diseases - comprising several peptide(s) consisting of V3 region peptide of envelope Gp., 120, e complex including carrier protein.
                                                                                                                                                                                                                                                                                                                                                           26-MAR-1992;
14-AUG-1992;
15-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-OCT-2003
25-MAR-2003
22-APR-1994
                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                 Okuda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  to neutralise HIV-1 in the treatment of HIV infection caused by different strains. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                   Disclosure; Page 3; 35pp; Japanese
                                                                                                                                                                                                                                                                                        WPI; 1993-320455/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                V3 region; HIV; envelope gp120; vaccine; human; humoral response; cellular immunity; carrier protein; human serum albumin; HSA; keyhole limpet haemocyanin; KLH; multiple antigen peptide.
                                                                                                                                                                                                                                                                                                                                       (TSDT-)
                                                                                                                                                                                                                                                                                                                                                                                                          19-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                  30-SEP-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        W09318791-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV gp120 V3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR41336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR41336 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
 15
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                                           Similarity
8; Conserv
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8; Conserv
                                                                                                                                                                                                                                                                                                                                       TSD
RAFVTIGK 22
                     RAFVTICK 8
                                                                                           25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RAFVTICK 8
                                                                                          ₽,
                                             Conservative
                                                                                                                                                                                                                                                                                                                                       줐
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               region peptide HIV-III B.
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                                                                                                                                                                                                                                                                                                                                                           92JP-00098602.
92JP-00237648.
93JP-00054239.
                                                                                                                                                                                                                                                                                                                                                                                                           93WO-JP000327
                                                     100.0%;
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Pred. No. 0.22;
0; Mismatches
                                                      Score 39;
Pred. No.
                                           Mismatches
                                                     DB 2;
0.22;
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                                           0,
                                                                 Length
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                                           Indels
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                                                                 25;
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                                          Gaps
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RESULT 148
ARA41330
ID ARA4133
XX ARA413:
XX ARA413:
XX 25-MAR
DT 25-MAR
DT 21-APR
XX 21-APR
XX HIV; h
KW enhanc
XX Human
XX WO9319
XX WO9319
XX WO9319
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PM WO915
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PM WOHL/
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X
ARASSETUT 149
ARASSES
XX ARASSES
AC ARASSES
XX 25-MAR
DT 06-SEP
XX Virus
XX Virus
XX Virus
XX Human
XX Synthe
XX Synthe
XX WO9308
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Best Local S
Matches 8
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                                                   Synthetic
                                                                                                  Human immunodeficiency virus;
                                                                                                                                                     Virus neutralising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The HIV gp120 epitope is used to isolate, create or evolve novel mols. including (in)organic and biomolecules such as proteins, peptides, nucleic acids, oligonucleocides, lipids, and polysaccharides for use as reactants, catalysts, enzymatic cofactors, repressors, enhancers, hormones and binders for a wide variety of substrates in industrial and therapeutic products. This epitope was isolated from variable region 3 of HIV gp120 (amino acids 271-295). (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                            25-MAR-2003
06-SEP-1993
                                                                                                                                                                                                                                                                                           AAR36587;
                                                                                                                                                                                                                                                                                                                                            AAR36587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 151; Page 147; 165pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Obtaining a novel mol. - capable of a desired interaction with a substrate of interest and a selection molecule expressed by the host.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1993-320737/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV; haemagglutinin; reactants; catalysts; cofactors; repreenhancers; hormones; binders; human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wohlstadter JN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-MAR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-SEP-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV gp120 epitope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9319170-A1
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21-APR-1994
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8; Conserv
                                                                                                                                                                                                                                                                                                                                         standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
ilarity 100.0%;
Conservative 0
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(first ent
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(first en
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                                                                                                                                                     epitope of
                                                                                                                                                                                                                                                                                                                                       peptide;
                                                                                                                                                                                                         entry)
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                                                                                               gp120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 39; DB 2; Pred. No. 0.22; Mismatches (
                                                                                                                                                   envelope glycoprotein of
                                                                                                                                                                                                                                                                                                                                         ₽
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                                                                                                  gp160; EGP; VNE; immunity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 25;
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WO9308836-A1

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RESULT 150
AAW72819
ID AAW728
XX AAW728
XX 17-OCT
DT 13-JAN
XX HIV-1;
XX HIV-1;
XX HIV-1;
XX Human
XX FF Peptid
FT Peptid
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Best Local :
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 29-MAY-1987;
24-DEC-1987;
25-APR-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-OCT-1991;
28-OCT-1991;
28-OCT-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A novel method of enhancing the immunogenicity of an envelope glycoprotein (EGP) of a virus (esp. HIV gpl20 or gpl60) in a host comprises admin. to the host at least one EGP of the virus in an amt. sufficient for priming vaccination and at least one peptide derived from an amino acid sequence of the EGP (e.g. the sequence shown), where the peptide comprises at least one virus-neutralisation epitope (NVB). The complex is able to enhance the induction of neutralising antibodies to the virus and to confer long lasting immunity, longer than 6 months. See also AAR36567-86. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                            HIV-1;
                                                                                                                                                                                                                                                                                                                                                        17-OCT-2003
13-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Enhancing immunogenicity of viral envelope glycoprotein - by co-
administration of viral envelope glycoprotein itself, and an oligopeptide
                                                               04-MAR-1993;
                                                                                                                                                                     Peptide
                                                                                                                                                                                                    Peptide
                                                                                                                                                                                                                                                 Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                          HIV-1; gp120; epitope; monoclonal antibody; envelope; neutralise; inhibit; infection; T-cell; inhibit syncytium formation; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                       AAW72819;
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW72819 standard; peptide; 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       derive.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-MAY-1993
                                                                                              10-NOV-1998.
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                                                                                                                         US5834599-A
                                                                                                                                                                                                                                                                                                                          gp120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 100.8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |||||||
RAFVTIGK 22
                                                                                                                                                                                                                                                                                                                          epitope 294 to 318.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 82; 107pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PASTEUR
                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
 87US-00057445.
87US-00137861.
89US-00343540.
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91US-00782241.
91US-00782252.
                                                              93US-00026276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92WO-EP002459
                                                                                                                                                                     /label= peptide_a
11. 25
                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                         /label=
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                                                                                                                                                       peptide_b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 39; DB 2
Pred. No. 0.22;
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The present invention describes an antibody conjugate comprising an antibody (Ab) which binds to an epitope within amino acid residue 308-322 of gp120 and neutralises HIV-1, conjugated with a cytotoxic agent, an anti-viral agent or an agent which facilitates passage through the blood brain barrier. Also described is an antibody conjugate as above but where the Ab binds to an epitope within amino acid residue 298-312 of gp12 which neutralises HIV-1. The present sequence represents an HIV-1 gp120 epitope corresponding to positions 294 to 318. The Ab are monoclonal Ab which bind to the gp120 protein on the envelope of HIV-1. They inhibit the infection of T-cells and also inhibit syncytium formation. The antibodies are group specific and neutralise different strains and isolates of HIV-1. The antibodies have a variety of uses, including the treatment and prevention of AIDS and AIDS related complex. They are especially used to kill infected T-cells. (Updated on 17-OCT-2003 to
Sequence 25 AA;
                                          standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibody conjugate comprising monoclonal antibody - which binds to epitope within amino acid residue of gp120 which neutralises HIV-1
                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Col 8; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                            conjugated with, e.g. cytotoxic agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-008810/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JUN-1992;
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                                          OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92US-00895197
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                              Best Local Similarity
Matches 8; Conserv
                                              Query Match
18
               1 RAFVTIGK 8
RAFVTIGK 25
                               100.0%; ilarity 100.0%; Conservative 0
                                0;
                                        Score 39;
Pred. No.
                                Mismatches
                                       0.22;
                                               DB 2;
                                0
                                Indels
                                0
                                Gaps
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Length 25;

0

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RESULT 151
AAW87618
ID AAW876
XX AAW876
AC Epitop
KW Epitop
KW Epitop
KW Epitop
KW Syncyt
KW AIDS-r
KW Viral
XX Viral
XX US5854
XX US5854
XX 29-MAY
PR 22-SEP
XX 29-MAY
PR 24-DEC
PR 26-SEF
XX A (TANO-
                                                                                                                                                                                                           17-OCT-2003
20-MAR-2003
                                                                                                                                                                                                                                      AAW87618;
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                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                           (revised)
                                                                                                                                                                                                                                                     peptide;
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Epitope; gp120 protein; monoclonal antibody; HIV-1; antibody BATi antibody BAT085; T cell infection inhibition; syncytia formation; acquired immune deficiency syndrome; AIDS; AIDS-related complex; passive immunisation; antiviral; cytotoxic; Human immunodeficiency virus 1.

Epitope of HIV-1 gp120 protein.

03-MAR-1999

(first entry)

US5854400-A.

22-SEP-1992; 29-DEC-1998. 9208-00950571

29-MAY-1987; 24-DEC-1987; 26-SEP-1991; 87US-00057445. 87US-00137861. 91US-00767533.

(TANO-) TANOX

WPI; Fung

1999-095002/08

MSC,

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RESULT 152
AAE09522
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                                                                                                                                                                                                                                                                                                                                                     Mucin; cytostatic; immunostimulant; cell mediated immune response; carcinoma; adenocarcinoma; breast cancer; dendritic cell; vaccine; gene therapy; CTL; cytotoxin T-lymphocyte.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antibodies are designated BAT123, 267 and 085. Monoclonal antibodies neutralise HIV-1, inhibiting both infection of T cells and formation of syncytia, so are used to treat acquired immune deficiency syndrome (AIDS) and AIDS-related complex, by passive immunisation, as carriers of cytotoxic or antiviral agents, and in extracorporeal systems. They can also be used as immunoassay reagents (for diagnosis or measurement of viral load) and to screen for neutralising epitopes, potentially useful in vaccine development. (Updated on 20-MAR-2003 to correct PR field.) (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                         01-FEB-2000; 2000AU-00005369.
14-JUN-2000; 2000US-00593870.
                                                                                                                                                                                                                                                                  09-AUG-2001.
                        Disclosure; Page 19;
                                                    Immunostimulant peptide, us an epitope of the non-VNTR,
                                                                                                  WPI; 2001-541537/60
                                                                                                                                                                                                                                     01-FEB-2001; 2001WO-AU000090
                                                                                                                                                                                                                                                                                                WO200157068-A1
                                                                                                                                                                                                                                                                                                                                                                                                                   Human
                                                                                                                                                                                                                                                                                                                                                                                                                                               19-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAE09522 standard; peptide; 25 AA.
                                                                                                                                                             (AUST-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bind
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents an immune deficiency virus (HIV)-1. The sequence represents an immune deficiency virus (HIV)-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Col
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monoclonal antibodies directed against regions of gpl20 of humar deficiency virus-1 - are neutralising and able to inhibit infect cells and formation of syncytia, used for treatment, prevention dispnosis of acquired immune deficiency syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present sequence represents an epitope of the gp120 protein of human ne deficiency virus (HIV)-1. The sequence comprises amino acids 298 22 of gp120. The specification describes monoclonal antibodies which to sequences derived from the present epitope. Specifically, these
                                                                                                                                                                                                                                                                                                                         immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                  immunodeficiency virus
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                                                                                                                                                             AUSTIN
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                                                                                                                                                             RES INST
                                                                                                                               Pietersz
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                       84pp; English
                                                                                                                              GA,
                                                                    used
                                                    ed as an anti-carcinoma non-leader region of a
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                                                                                                                              Apostolopoulous
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Pred. No. 0.22;
; Mismatches (
                                                   region
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inhibit infection of '
                                                                   vaccine,
                                                      mucin
                                                                 comprises
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The patent

discloses

peptide or polypeptides capable

of eliciting

an

frame comprising a first sequence, where the first sequence encodes one or more segments of tumour-associated antigen SSX-2, which comprises a sequence of 188 amino acids (SEQ ID NO: 40), where the first sequence does not encode the complete SSX-2 antigen, and where each segment comprises an epitope cluster, the cluster comprising or encoding at least two amino acid sequences having a known or predicted affairly for a same MHC receptor peptide binding cleft. Also described are: an isolated

sequence

The invention

describes an isolated nucleic acid (I) comprising

þ

reading

260pp; English.

Disclosure; SEQ ID NO 131;

Novel nucleic acid WPI; 2004-517003/49.

inducing

an

immune

response encoding

and

in treating

cancer.

tumor-associated antigen

SSX-2, useful in

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ADQ10566
ID ADQ10566
ID ADQ105
AC ADQ105
XX ADQ105
XX ID Human
XX Immuno
XW SSX-2
XW Epitop
XW Cepitop
XW Cytoto
XX OYOOT
XX IMMUNO
XW SX-2
XW IMMUNO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunostimulant; cytostatic; vaccine; tumour-associated antigen SSX-2; SSX-2 antigen; epitope cluster; MHC receptor peptide binding cleft; immunogenic composition; immune response; cancer; vaccine vector; epitope liberation; human leukocyte antigen; HLA A2-specific CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Simard JJL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SIMA/)
(DIAM/)
(QIUZ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-NOV-2001; 2001US-0336968P.
07-NOV-2002; 2002US-00292413.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-FEB-2004; 2004US-00777053.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2004132088-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytotoxic T lymphocyte; T-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus T-cell epitope seqid 131.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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8; Conserv
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Pred. No.
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RESULT 154
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Best Local S
Matches 7
bond, thus giving the peptide a loop-like configuration, which is utilised to enhance the immunogenic properties of the peptide. The peptide is capable of eliciting, and/or binding with, neutralising antibodies. The neutralising domain is bounded by cysteine residues which occur at positions 296 and 331. Peptides can be used as immmunogens or screening reagents to generate or identify poly- or MAbs. See also AAR0427-R04506 and AAR04273-Q04279. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                           Peptide 135 comprises segments of the Principal Neutralising Domain (envelope protein) from isolate HIV-IIIB. The last Cys resisue is added for the purpose of crosslinking to carrier proteins. Cysteine residues can be added so that that residues at or near both ends form a disulfide
                                                                                                                                                                                                                                                                                                                                                                                                                        Principal neutralising domain of HIV variants - used peptide(s) and antibodies for diagnosis, prophylaxis therapy therapy of HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               frame; and an immunogenic composition comprising (I) or the polypeptide of (I). (I) is a nucleic acid encoding a tumour-associated antigen SSX-2 comprising a fully defined sequence of 188 amino acids (SEQ ID NO: 40). The nucleic acid, the encoded antigen, and composition are useful in inducing an immune response and in treating cancer. Expression cassettes are used in vaccine vectors. This is the amino acid sequence of a T-cell epitope MHC ligand associated with methods, therapies and compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1990-147824/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-OCT-1988;
01-JUN-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-OCT-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV-IIIB; peptide 135; principal neutralising domain; antibodies,
diagnosis; prophylaxis; therapy; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-SEP-2004
25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus peptide 135
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9003984-A.
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(revised)
(first entry)
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89US-00359543.
89US-00407663.
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                                                                                                                                                                                                                                                                                                                                                                      108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Javaherian K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 37; DB 8;
Pred. No. 1.8e+06;
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맑 5

RAFVTIGO 15 RAFVTIGK 8 Best Loc Matches

Local Similarity hes 7; Conserv

Conservative

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Indels

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Gaps

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ARESULT 155
ARAG6430
ID ARAG643
XX ARAG643
XX ARAG643
XX 25-MAR
DT 25-MAR
DT 25-MAR
DT 3-AUG
XX T cell
XX T cell
XX Cluste
XX Synthe
XX W09426
XX 24-NOV
XX 24-NOV
XX 14-MAY
XX (USSH
XX WP; 13-MAY
XX T respon
PT for th
XX PR HYP; 1
XX PR HYP; 1
XX PR HYP; 13-MAY
XX (USSH
XX WP; 11-MAY
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                                                                                                                                                 Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-R66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups c sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-15, the Lys residue at position 15 in peptide 18 has been replaced by a Gln residue. (Updated c25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polypeptide inducing helper T cell, cytotoxic T cell and responses - to target antigen in hosts of different MHC | for therapeutic or prophylactic vaccines against HIV.
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03-AUG-1995
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                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-006707/01.
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(first entry)
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87.5%;
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Score 35; DB Pred. No. 1; 1; Mismatches
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Pred. No. 0.63
1; Mismatches
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RESULT 157
AAW22329
ID AAW223
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AC AAW223
XX
T7-OCT
DT 18-SEP
XX
DE HIV-1
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AAR66424
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Matches
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HIV-1 clinical strain 9622 gp120 V3 loop peptide
                            17-OCT-2003
18-SEP-1997
                                                                        AAW22329;
                                                                                              AAW22329 standard; peptide;
                                                                                                                                                                                                                                                                                  Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polypeptide inducing helper T cell, cytotoxic T cell and antibodies responses - to target antigen in hosts of different MHC haplotypes, esp. for therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-MAY-1993;
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03-AUG-1995
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                                                                                                                                                                                                                           Similarity 7; Conserv
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K
                            (revised)
(first en
                                                                                                                                                                                                                           Conservative
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(first entry)
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                            entry)
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                                                                                                                                                                                                                                       89.7%;
                                                                                                 19
                                                                                                                                                                                                                       Score 35; DB Pred. No. 1; 0; Mismatches
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                                                                                                                                                                                                                       Gaps
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The invention relates to a novel monoclonal antibody (MAb) NM03 which CC binds to epitopes from the human immunodeficiency virus type 1 (HIV-1). CC The antibody was raised conventionally by immunising Balb/c mice with CC purified live HIV-1 MN, then isolating splenocytes and fusing them to p3-CC X63-Ag8-U1 cells. Hybridomas were then screened with membrane fractions CC from infected and non-infected H9 cells. The MAb was observed to bind to CC a protein band of 120 kD on a Western blot of separated, denatured HIV-1 CC proteins. This binding was shown to be between residues 320-327 by CC epitope mapping by ELISA and competitive binding. The ability of the MAb CC inhibit infection of cells by HIV-1 shown by infecting H9 cells with CC live strains of HIV-1 and testing infection by a p24 assay. This peptide CC sequence represents the V3 loop region from HIV-1 clinical strain 9622, CC where the MAb NMO3 binds. The MAb can be used for the passive CC immunisation of humans susceptible to, or infected with HIV-1. (Updated CC con 17-CCT-2003 to standardise CS field)
Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                Example 3; Col 10; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                          Monoclonal antibody specific for human immunodeficiency virus type 1 MN strain - for passive immunisation against infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-APR-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yoneda Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NISP ) NISSIN SHOKUHIN KAISHA LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ohno T,
AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                        immunisation against infection.
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               Matches
                       Query Match
Best Local
              Similarity 7; Conserv
RAFVTICK 8
               Conservative
                      89.7%;
              Score 35; DB
Pred. No. 1.3
0; Mismatches
                    DB
1.3;
                             2
                             Length 19
              0,
              Gaps
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0,

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RESULT 158
AAW62892
ID AAW628
XX AAW628
XX 30-SEP
XX 30-SEP
XX Wonocl
KW HIV st
XX Wonocl
KW HIV st
XX Synthe
XX JP1018
XX JP1018
XX PF 25-DE(
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AAW62892;

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RTFVTIGK 13

AAW62892 standard; peptide; 19

30-SEP-1998 (first entry)

Monoclonal antibody; HIV-1gp120; gp160; infection; H9 cell; HIV strain MN; treatment; human HIV infection. Peptide sequence of the specification.

Synthetic.

JP10182489-A

07-JUL-1998.

25-DEC-1996; 96JP-00344904

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RESULT 159
ABB05775
ID ABB057
XX ABB057
XX ABB057
XX POLYFU
RW POLYFU
KW POLYFU
KW POLYFU
KW ARTIFI
XX WO2001
XX POLYFU
PT CULTUR
XX WPI; Z
XX WPI
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Best Local
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The present invention describes a polyfunctional base sequence (N1) having two or more functions in different reading frames. Also described are: (1) a method for producing N1 and artificial gene expression vectors comprising N1; (2) transgenic non-human animals comprising N1; and (3) treatments and diagnostic reagents containing an artificial protein, artificial tissues or high molecular weight artificial proteins. N1 is useful for creating industrially useful artificial matrix proteins for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW62889-900 represent peptides used to identify a peptide sequence (AAW62874) present in HIV-1gp120 or gp160 which is bound by the monoclonal antibody of the invention. The antibody neutralises in the infection of H9 cell by an active HIV strain MN according to the analytical method. The antibody is used for treatment of human HIV
                                                                                                                                                                                                                                                                                                           Polyfunctional base sequence having two or more functions in different reading frames, useful for producing artificial matrix proteins for cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polyfunctional base sequence; microgene; industrial; cell artificial matrix protein; transgenic animal; HIV.
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07-MAY-2002
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                                                                                                                                                                                                                            Example 1;
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-098069/13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABB05775;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NISC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JUN-2001; 2001WO-JP005116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200196558-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NISP ) NISSIN SHOKUHIN KAISHA LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RTFVTIGK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     related
                                                                                                                                                                                                                            Page 46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ş
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide;
                                                                                                                                                                                                                         61pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             89.7%;
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Pred. No. 1.3;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2
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RESULT 160
RAD15657
ID AAO156
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Best Local (
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Best Local (
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                                                                                                                                                                                                                                            The invention comprises an artificial protein which induces a strong immune response to a peptide group (the protein contains all or part of the peptide group). The artificial protein assists the formation of high order protein structure and/or assists the antigen presentation of immunocompetent cells. The artificial protein of the invention is useful for inducing a strong immune response and the preparation of effective antibodies to specific antigens, especially HIV. The present amino acid sequence represents a peptide that was used in the invention
                                                                                                                                                                                                    Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Artificial protein capable of inducing a strong immune response to peptide group for assisting antibody production in vivo to viruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shiba K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Strong immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell culture. The present sequence represents a peptide which is used an example from the present invention. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 6; Page 5; 77pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     other antigens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-519151/55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antigen presentation; HIV.
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                                               1 RAFVTICK 8
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                                                                                                  Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
7; Conserve
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  RTFVTICK 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ohno
                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                         89.7%;
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87.5%;
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                                                                                                                         Score 35;
Pred. No.
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ₽
                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                    Length 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in vivo to viruses and
                                                                                                    Indels
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RESULT 161 AAR62151 ID AAR62:

AAR62151 standard; peptide; 8

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RESULT 162
AAR62138
ID AAR621
XX AAR621
XC AAR621
XX 27-AUG
DT 27-AUG
DT 25-MAY
XX U2-MAY
XX XX Small
KW Small
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Best Local S
Matches 7
Small ribonucleoprotein complex; UI snRNP; 70K protein; epitope; autoantibody; immunoinfective cluster virus; nuclear protein anti; systemic rheumatic disorder; human immunodeficiency virus; HIV-1; systemic lupus erythematosus; mixed connective tissue disease; systemic lupus erythematosus; mixed connective tissue disease;
                                                                                                                       27-AUG-2003
25-MAR-2003
02-MAY-1995
                                                                                                                                                                                AAR62138;
                                                                                                                                                                                                                                                                                                                                                                                                                   The UI snRNP is the target of high-titre, high avidity autoantibodies occurring in the systemic rheumatoid disorders of mixed connective tissue disease, scleroderma and systemic lupus erythematosus. It has been found that some sites in the UI snRNP 70K protery (see AARG2120-R62135) are homologous to sites in HIV-1 gpl120/41 (AARG2136-R62152) and that anti-RNP autoantibodies can be used to neutralise HIV-1. (Updated on 25-MAR-2003 to correct DN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-AUG-2003
25-MAR-2003
02-MAY-1995
                                                                                                                                                                                                            AAR62138
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 56; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   patients
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Douvas A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9420141-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoantibody; immunoinfective cluster virus; nuclear protein anti-
systemic rheumatic disorder; human immunodeficiency virus; HIV-1;
systemic lupus erythematosus; mixed connective tissue disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Small ribonucleoprotein complex; UI snRNP; 70K protein; epitope; autoantibody; immunoinfective cluster virus; nuclear protein ant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYSC-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR62151;
                                                                                           gp120/41 protein motif similar to UI
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                                                                                                                                                                                                                                                                                                                                     Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for treating immunoinfective cluster virus infections - ies or fragments characteristic of auto antibodies produc with rheumatic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              UNIV SOUTHERN
                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                         RAFVTIG 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Takehana Y,
                                                                                                                                                                                                                                                                                                                                  87.2%; Score 34; DB llarity 100.0%; Pred. No. 1.: Conservative 0; Mismatches
                                                                                                                                      (revised)
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(revised)
(first en
                                                                                                                       (first entry)
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                                                                                                                                                                                                         peptide;
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o. 1.8e+06;
0;
                                                                                        snRNP 70K protein.
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                                            antigen;
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RESULT 163
ADK68768
ID ADK687
XX ADK687
XX ADK687
XX Epitop
XX Epitop
XX Epitop
XX Epitop
XX Virus;
OS Human
XX US2003
XX US2003
XX I1-DEC
XX PF 07-NOV
XX PF 07-NOV
XX PF 07-NOV
XX PA (SIMA, PA (DIAM, PA (DIAM, PA (LEIX, PI (LEIX, PA (LEIX, PI (LEIX, PA (L
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Best Local S
Matches 7
                                                                           (SIMA/)
(DIAM/)
(QIUZ/)
                                                                                                                                                                                   07-NOV-2001;
                                                                                                                                                                                                                                      07-NOV-2002; 2002US-00292413.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease, scieroderma and systemic lupus erythematosus. It has been found that some sites in the UI snRNP 70K protein (see AAR62120-R62135) are homologous to sites in HIV-1 spi20/41 (AAR62136-R62152) and that anti-RNP autoantibodies can be used to neutralise HIV-1. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                              11-DEC-2003
                                                                                                                                                                                                                                                                                                                                                US2003228634-A1
                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protozoacide; fungicide; T-ce
cytotoxic T lymphocyte; CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    epitope liberation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Epitope liberation-related peptide SeqID131.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK68768;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADK68768 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Methods for treating immunoinfective cluster virus infections antibodies or fragments characteristic of auto antibodies procepatients with rheumatic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The U1 snRNP is the target of high-titre, high avidity autoantibodies occurring in the systemic rheumatoid disorders of mixed connective ti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 52; 106pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus
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                                                  SIMARD J J L.
DIAMOND D C.
QIU Z.
LEI X.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                      2001US-0336968P
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                                                                                                                                                                                                                                                                                                                                                                                                                                       n; substrate; proteasome; cytostatic; antibacterial; gicide; T-cell activator; vaccine; housekeeping epitope; ocyte; CTL; adoptive immunotherapy; neoplastic cell; protozoan; fungus; housekeeping proteasome system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Y,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              core 34; DB 2; L
Pred. No. 1:8e+06;
Mismatches 0;
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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                               27-AUG-2003
25-MAR-2003
03-MAY-1995
                                                                                                                                                                                                                                                    epitope; autoantibody; immunoinfective cluster virus; nuclear protein antigen; systemic rheumatic disorder; human immunodeficiency virus; HIV-1; systemic lupus erythematosus; mixed connective tissue disease; scleroderma; glycoprotein 120.
                                                                                                                                                                                                                                                                                                                                                          HIV-1 gp120 V3 loop neutralising domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR62165;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR62165 standard; peptide; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying polypeptide suitable for epitope e.g., housekeeping epitope liberation by contacting substrate polypeptide comprising epitope of interest, with proteasome, and assaying for liberation of epitope.
                                                          11-MAR-1993;
                                                                                                                                                                            WO9420141-A1
                                                                                                                                                                                                                   Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 131; 67pp; English
                  (UYSC-) UNIV SOUTHERN CALIFORNIA
                                                                                               10-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-167209/16
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                                                          93US-00029850.
                                                                                               94WO-US002631.
                                                                                                                                                                                                                                                                                                                                                                                                 entry)
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Pred. No. 1.8e+06;
0; Mismatches 1
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RESULT 165
AAW54661
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Previous immunological analyses of the V3 loop of HIV-1 (AAR62159) have localised the main neutralising domains. The target of more than 80% of neutralising antibodies in HIV-1 infected sera from AIDS patients has now been found to overly the consensus binding sequence and domain A epitopes of the U1 snRNP 70K protein. In AIDS, antibody titres are too low to arrest the disease; however, the homologous sequences in 70K are immunodominant targets of autoantibodies in the systemic rheumatoid disorder of mixed connective tissue disease. The titers of such autoantibodies exceed 10 power 7. The anti-snRNP autoantibodies will cross-react with HIV-1 epitopes and are useful for treating HIV infection. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-ANG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Methods for treating immunoinfective cluster virus infections - utilise antibodies or fragments characteristic of auto antibodies produced by patients with rheumatic disorders.
                                                                                       Increasing uptake and presentation of antigen(s) - residue(s) to antigen for increasing T cell respons vaccines against viral infection(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                   25-SEP-1998
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                                                               Disclosure; Page 29; 47pp;
                                                                                                                                                                       Koning
                                                                                                                                                                                                                            26-SEP-1996;
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                                                                                                                                                                                                                                                                                                             WO9813378-A1
                                                                                                                                                                                                                                                                                                                                                                  vaccine; treatment.
                                                                                                                                                                                                                                                                                                                                                                            Mannose; antigen; antigen-presenting cell; mannosylated peptide; T cell;
                                                                                                                                                                                                                                                                                                                                                                                                        Peptide from HIV-1 gp120 314-322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW54661 standard; peptide;
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                                                               English.
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Pred. No.
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                                                                                                   en(s) - by adding mannose response, useful in, e.g.
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The peptides AAM54559-W54809 are examples of peptides to which at least (preferably 2) mannose can be attached to increase their uptake as antigens by antigen-presenting cells. Uptake of agonist mannosylated peptides will increase the T cell response, whereas uptake of antagonist

antagonist

888888**%**&

Sequence 10

\$

DB 2;

Length 10;

parasites

peptides blocks the T cell response. Blocking binding of immunogenic autoantigens can be used in treatment of type I diabetes, rheumatoid arthritis, graft rejection etc., also to induce T-cell non-responsiveness. Vaccines containing mannosylated antigen are used to prevent or treat infections by, e.g. bacteria, viruses, fungi, helmir

helminths

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RESULT 166
AAR62167
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                    Previous immunological analyses of the V3 loop of HIV-1 (AAR62159) have localised the main neutralising domains. The target of more than 80% of neutralising antibodies in HIV-1 infected sera from AIDS patients has now been found to overly the consensus binding sequence and domain A epitopes of the U1 snRNP 70K protein. In AIDS, antibody titres are too low to arrest the disease; however, the homologous sequences in 70K are immunodominant targets of autoantibodies in the systemic rheumatoid disorder of mixed connective tissue disease. The titers of such autoantibodies exceed 10 power 7. The anti-snRNP autoantibodies will cross-react with HIV-1 epitopes and are useful for treating HIV infection. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 27-AMR-2003 to correct PN field.)
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25-MAR-2003
03-MAY-1995
                                                                                                                                                                                                                                                                     Methods for tr
antibodies or
                                                                                                                                                                                                                                                                                                                                                  Douvas A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nuclear protein antigen; systemic rheumatic disorder; nuclear protein antigen; systemic rheumatic dupus erythematosus; human immunodeficiency virus; HIV-1; systemic lupus erythematosus; mixed connective tissue disease; scleroderma; glycoprotein 120; Ul snall nuclear ribonucleoprotein; Ul snRNP 70K protein.
                                                                                                                                                                                                                                                                                                                       WPI; 1994-302689/37
                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                               (UYSC-) UNIV SOUTHERN CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           epitope; autoantibody; immunoinfective cluster virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1 gp120 V3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR62167;
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      loop domain containing U1 snRNP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide;
                                                                                                                                                                                                                         106pp; English
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RESULT 167
AAW76852
ID WAM768
XX AAW768
XX AAW768
XX AAW768
XX Pusion
XX B cell
XX microb
XX mic
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus sp.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 10; Page 119; 154pp; English
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100.0%;
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Pred. No.
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This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl10 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, so prevents or inhibits 'bystander' apoptosis of unifected T cells that are bound to contact manning to that the contact in maximizing induction of antibodies.

are bound to gp120 protein, maximising T cell immunity

induction of protective

밁 δ Matches Query Match Best Local Sequence 11 AA; s ۲ Similarity 7; Conserv RAFVTIG 7 Conservative 87.2%; 500 100.0%; Pr Score 34; Pred. No. Mismatches 1. B 2 0, Length 11 Indels ٥, Gaps

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Sequence 11 AA;

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                                                                                                                                                                                               neuroprotectant, virucidal, bactericidal, antiparasitic, fungicidal and cytostatic activities. (I) are useful, in human or veterinary medicine, in pharmaceutical compositions (for treating immune disorders, e.g. immune deficiency, autoimmune states, hypersensitivity, allergy, graft rejection, infection, hormonal disorders and central nervous system diseases), also, where (I) is a MHC ligand (Ia), in vaccines for treatment or prevention of: (i) viral, bacterial, parasitic or fungal infections; or (ii) of cancers. A particular application is in antimelanoma vaccines. (I) are also useful for in vitro diagnosis of diseases associated with interactions between MHC and (I), e.g. melanoma and human immunodeficiency virus infection. AAM98898 to AAM99592 represent peptides which can be used in pharmaceutical compounds from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a pharmaceutical compound (I) that contains an N-terminal glutamic acid (Glu) or glutamine (Gln) residue in the form of an addition salt with a strong, physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition containing at least one (I); (b) a vaccine containing at least one (I) where this is a major histocompatibility complex (MHC) ligand (Ia); (c) method for in vitro diagnosis of diseases associated with the presence (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process for preparing (I). (I) has immunomodulator, endocrine, antiallergic, neuroprotectant, virucidal, bactericidal, antiparasitic, fungicidal and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stabilized pharmaceutical containing N-terminal glutamic acid or glutamine, useful e.g. in anti-melanoma vaccines, is an addition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunomodulator; antiallergic; endocrine; neuroprotectant; virubactericidal; antiparasitic; fungicidal; cytostatic; medicine; pharmaceutical; immune disorder; immune deficiency; autoimmune;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pharmaceutical; immune disorder; immune deficiency; autoimmune; hypersensitivity; allergy; graft rejection; infection; hormonal disorder central nervous system disease; cancer; melanoma; anti-melanoma vaccine;
                                                                                                                                                               Sequence 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Klinguer-Hamour C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-MAR-2001; 2001WO-FR000872.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Glutamic acid; glutamine; vaccine; major histocompatibility complex; MHC
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                                                                                Local Similarity
les 7; Conserv
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                                        RAFVTIG 7
                                                                                87.2%; Score 34; DB ilarity 100.0%; Pred. No. 1. Conservative 0; Mismatches
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                                                                                                   DB 4;
1.4;
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                                                                                                                       Length 12;
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Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                           The peptides of the invention comprise the beta-turn AA SQ GPG or GPGR at positions 312-314 or 312-315 in the AA numbering of HTLV-IIIB (BH10) and flanking AA SQs having a length equal to or greater than 1 and pref. equal to or greater than 2 AAs; variants in which the GPG or GPGR SQ has been replaced by a different beta-turn SQ; and variants in which the free NH2-terminal gp AA and/or the free carboxyl terminal gp. AA has been blocked or modified otherwise. (Updated on 27-AUG-2003 to correct OS field.)
                                                                                                                                                       Sequence 15 AA;
AAR33460 standard; peptide; 15
                                                                                                                                                                                                                                                                                       Disclosure; Page 3; 7pp; English.
                                                                                                                                                                                                                                                                                                              Oligopeptide(s) corresp. to beta-turn variable region of gp.120 - for diagnosis of and prodn of vaccines against AIDS and ARC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-OCT-1988;
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30-MAR-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      causing or related virus.
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(UNAM ) UNIV VAN AMSTERDAM.
(UYAM-) UNIV AMSTERDAM ZIEKENHUI.
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                                                                                  RAFVTIG 7
                                                                                                           Conservative
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(first entry)
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                                                                                                                    87.2%;
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                                                                                                                    Score 34; DB 1;
Pred. No. 1.7;
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                                                                                                                                Length 15;
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25-MAR-2003
17-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                           produce a peptide which is more active in eliciting a cytofoxic T lymphocyte (CTL) response. It can be used for the treatment and/or prophylaxis of HIV infection. (Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent clashes with ongoing US granted patent numbers. For further information please visit the Derwent web site at www.derwent.com/dwpi/updates/ntis us.html.) (Updated on 25-MAR-2003 to correct PF field.) (Updated on 27-AUG-2003 to correct OS field.)
           epitope; autoantibody; immunoinfective cluster virus; nuclear protein antigen; systemic rheumatic disorder; human immunodeficiency virus; HIV-1; systemic lupus erythematosus; mixed connective tissue disease; scleroderma; glycoprotein 120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The peptide corresp. to residues 319-329 of HIV-1 strain IIIb gp. envelope glycoprotein. It is activated by cleavage with a protease produce a peptide which is more active in eliciting a cytotoxic T
                                                                                                                                                                      AAR62166;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-MAR-1992;
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                                                                                   HIV-1 gp120
                                                                                                                                                                                                  AAR62166 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide(s) for stimulation of cytotoxic T cells specific for HIV-1 which correspond to residues 318-327 of HIV-1 gp 160 envelope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berzofsky JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JAN-1988;
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                                                                                  loop dominant neutralising domain in chimpanzees
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87.5%;
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Pred. No. 1.7;
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Matches
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                                                                                                                                                                          T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                  HIV-1 IIIB peptide 18-12.
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03-AUG-1995
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                                        14-MAY-1993;
                                                                   13-MAY-1994;
                                                                                              24-NOV-1994
                                                                                                                         WO9426785-A1
                                                                                                                                                                                                                                                                                                                             AAR66427 standard; peptide; 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Methods for treating immunoinfective cluster virus infections - utilise
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    87.2%; Score 34; DB ilarity 100.0%; Pred. No. 1. Conservative 0; Mismatches
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                                                                                                                                                                                                                                                           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fragments characteristic of auto antibodies produced by rheumatic disorders.
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Berzofsky JA, Ahlers JD,

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                                                                                                  Example 1;
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                                                                                                                                                                                                                                                                                                                                                                       JA,
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                                                                                                                                                              inducing helper T cell, cytotoxic T cell and antibodies to target antigen in hosts of different MHC haplotypes, utic or prophylactic vaccines against HIV.
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Pred. No. 1.7;
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residues from the HIV-1 RF sequence (AAR66415) replaced residues HIV-1 IIIB sequence (AAR66414) to test the effect of each residu

Sequence 15 AA

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RESULT 174
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                                                                                                   Sequence 15
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03-AUG-1995
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glycoprotein gp120;
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RESULT 176 AAR33236

EP311219-A HTLV-IIIB.

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RESULT 177
AAP95348
ID AAP953
XX Uariab
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Best Local (
                                                                                                                                                                                                                                                                                                               Matches
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                                                              Vaccine; AIDS;
                                                                                       Variable region V3 sequence found in the envelope protein gp120 of an AIDS or ARC causing or related virus.
                                                                                                                              27-AUG-2003
30-MAR-1992
                                                                                                                                                                    AAP95348;
                                                                                                                                                                                             AAP95348 standard;
                                                                                                                                                                                                                                                                                                                                                                Sequence 16 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synergistic compsn. for treating HIV-1 infection - comprises antibody V3 loop of GP120 and antibody to CD4 binding site of GP120 or soluble
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20-JUL-1992;
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13-JUL-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV-1; human immunodeficiency virus; competition assay; AIDS; infection; CD4 binding site; soluble CD4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV-IIIB gp120 V3 loop epitope peptide RP135
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(first entry)
                                                              ARC; HIV; diagnosis
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92US-00916542
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                                                                                                                              entry)
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100.0%; Pred. No.
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(UNAM)
(UYAM-)
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Oligopeptide(s) corresp. to beta-turn variable region of gp.120 for diagnosis of and prodn of vaccines against AIDS and ARC.
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30-MAR-1992
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                                                                                                                                                                                                                                                     09-OCT-1987;
                                                                                                                                                                                                                                                                                                  07-OCT-1988;
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                                                                                                               Goudsmit J,
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(UNAM ) UNIV VAN AMSTERDAM.
(UYAM-) UNIV AMSTERDAM ZIEKE
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7; Conserve
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UNIV AMSTERDAM ZIEKE
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AMSTERDAM ZIEKENHUI.
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RESULT 179
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                                 The peptides of the invention comprise the beta-turn AA SQ GPG or GPGR positions 312-314 or 312-315 in the AA numbering of HTLV-IIIB (BH10) an flanking AA SQs having a length equal to or greater than 1 and pref. equal to or greater than 2 AAs; variants in which the GPG or GPGR SQ ha been replaced by a different beta-turn SQ; and variants in which the fr NH2-terminal gp AA and/or the free carboxyl terminal gp. AA has been blocked or modified otherwise. (Updated on 27-AUG-2003 to correct OS
Sequence 17
                                                                                                                                 Disclosure; Page 3; 7pp; English.
                                                                                                                                                      Oligopeptide(s) corresp. to beta-turn variable region of gp.120 for diagnosis of and prodn of vaccines against AIDS and ARC.
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30-MAR-1992
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Pred. No.
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Query Match Best Local Similarity

87.2%; 100.0%;

Pred.

34; DB

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Length 17;

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RESULT 180
AAR29241
ID AAR292
XX AAR292
XX AAR292
XX 24-OCT
DT 25-MAR
DT 14-APR
XX V3 loo
XW V3 loo
XW homolo
XX Human
XX Homan
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RESULT 181
AAR32407
ID AAR324
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Best Local S
Matches 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequences given in AAR29237-43 represent a portion of the V3 loop region of the gp120 envelope protein of various HIV isolates. These sequences can be used to define the specific isolate. All these viral varients exhibit complete homology at residues 7-11 of the given sequence and at least 36% homology with the remaining 12 amino acids of the sequence. Viruses containing these sequences are recognised by the heteroconjugate enzyme of the invention. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 24-OCT-2003 to standardise OS field)
                           25-MAR-2003
04-JUL-1993
                                                                                                          AAR32407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hetero-conjugate antibodies for treating HIV infections - comprise an antibody specific for an effector cell surface antigen and an antibody to a V3 loop of GP-L20 envelope protein of HIV.
                                                                                                                                                           AAR32407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 32; 69pp; English
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08-JUN-1994; 22-DEC-1994.

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Human immunodeficiency virus 1.

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25-MAR-2003
06-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nine peptides from the V3 regions of HIV-1 isolates IIIb, RF and MN were incorporated into tetravalent multiple antigen peptide systems (MAPS) (see AAR32406-14). Parallel groups of three peptides with chain lengths spanning from 11-24 residues were synthesised in MAPS formate for each isolate. ELIS assays demonstrated that antisera titers in mice were closely related to the length of the IIIb peptide used for the immunisation - the longer the stronger the response. There was no substantial antibody prodn. in mice against the other two series of peptides, RF (B4-B6), and MN (B7-B9), except for a low reactivity in the gp. immunised with B8 (MN isolate). Specificity tests of the B cell response demonstrated that the T cell epitope (AAR32415) also serves as a B cell epitope. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                        T-cell; epitope; HIV-1; core protein; p24E; B-cell; antigen; gp160; gag; pol; vaccine; multimeric peptide; AIDS; 3D organisation.
                                                                                                                                                                                                                                                                                                               T cell epitope derived from V3 isolate LAI.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel tandem synthetic HIV-1 peptide(s) - comprising T-cell epitope of gag protein linked to B-cell epitope of V3 loop protein of an HIV-I isolate.
                        Tandem synthetic HIV peptide(s) useful as immunogens - comprising protein T-cell epitope linked to env protein B-cell epitope.
                                                                                                                                                                    09-JUN-1994;
                                                                                                                                                                                                                       US5639854-A.
                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                       HIV; human immunodeficiency virus; gag; T-cell; B-cell;
V3 loop; vaccine; determinant; chimaeric.
                                                                                                                                                                                                                                                                                                              HIV B-cell strain LAI env protein V3 loop peptide
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20-OCT-1997
Disclosure; Col 21; 41pp; English
                                                                                                                                            09-JUN-1993;
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                                                                         The invention relates to new synthetic peptides comprising at least one amino acid sequence comprising an HIV gag protein T-cell epitope linked at its C- or N-terminus to an amino acid sequence comprising a B-cell epitope of the V3 loop of an HIV env protein, which can be used to generate vaccines against HIV-1. The T-cell epitope sequence is pref. selected from the T-helper determinant core peptides P24B, P24N, P24N, P24N and P24H while the B-cell epitopes are derived from HIV strains including CTLB-56, V3NN, CTLB-29, CTLB-55, SF2, LAI, IIIB, RF, Z6, 2054, 1714 and BXOB. The peptides are chimaeric and can be linked to a branched by backbone. This sequence represents the B-cell env protein V3 loop peptide from HIV-1 strain LAI. (Updated on 25-MAR-2003 to correct PF
Sequence 17
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                            Query Match
Best Local (
                    Matches
 1 RAFVTIG 7
                   Similarity 7; Conserv
                    Conservative
                            87.2%;
100.0%;
                    Score 34; DB; Pred. No. 2; 0; Mismatches
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Gaps

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RAFVTIG 17

AAW76848 standard; peptide; 17 ₿

AAW76848

25-JAN-1999 (first entry)

Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #18.

B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.

Mus sp. Homo sapiens

WO9836087-A1

20-AUG-1998

13-FEB-1998; 98WO-US002766.

13-FEB-1997; 97US-0040581P.

(AMNA-) AMERICAN NAT RED CROSS

'n Zambidis E

WPI; 1998-506315/43.

induce tolerance to the epitopes virus infection. New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency

Claim 10; Page 119; 154pp; English.

RESULT 184
AAW76848
ID AAW768
XX AAW768
XX AAW768
XX Pusion
XX E cell
KW Inman
KW prophy
KW Inman
KW prophy
KW Inman
KW prophy
KW INMA
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KW INMA
DE COLAIM
KW INMA
CC IMMUNG
CC OPATION
This sequence is an epitope used in the construction of a novel fusion immunoglobulin heavy chain (IgH) protein with a mammalian, especially human, IgH chain fused in frame at its N-terminus to one or more human immune deficiency virus (HIV) gpl20 epitopes. Fusion Ig proteins and/or transfected cells are used to tolerise subjects to gpl20 epitopes and to maintain this tolerance, particularly for treatment of HIV infection, optionally together with other therapeutic/prophylactic agents such as vaccines, chemotherapeutic agents and immune response modifiers. Such

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Best Local S
Matches 7
                                                                                Query Match
Best Local :
                                                                                                                                                          The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-cell epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitopes are generally designed based on the p24 core protein and the B-cell epitopes from the V3 loop of the gp120 protein from various HIV-1 strains. This peptide represents the V3 loop epitope from the HIV-1 strains. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                               Sequence 17
                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                   Synthetic human immunodeficiency virus-1 peptide(s) - contain epitope and B-cell epitope(s) are candidate vaccines against
                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-556461/47
                                                                                                                                                                                                                                                                                                                                                                                                                         Chong P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-JUN-1993;
09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US5817754-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunogen; vaccine; HIV-1; T-cell; B-cell; epitope;
V3 loop.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-1 strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-OCT-2003
25-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    proteins can be used against other diseases where an immune response is deleterious, e.g. microbial infection, tumours or autoimmune disease. Induction of tolerance suppresses production of antibodies against gpl20, so prevents or inhibits 'bystander' apoptosis of uninfected T cells that are bound to gpl20 protein, maximising induction of protective antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CONN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW67350;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW67350 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           und to gp120 protein, immunity
                                                                Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                         CONNAUGHT LAB
RAFVTIG 17
                                RAFVTIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RAFVIIG 7
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                                                                                                                                                                                                                                                                                                                                                                                                                         Klein MH,
                                                             87.2%; Sillarity 100.0%; Conservative 0;
                                                                                                                                  Ą,
                                                                                                                                                                                                                                                                                                  Col 21; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ДĄ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LAI gp120 V3 loop epitope peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93US-00073378.
94US-00257528.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-00464329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                         Sia
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                                                                                                                                                                                                                                                                                                                                                                                                                         CDY;
                                                           Score 34; DB;
; Pred. No. 2;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2;
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                                                              0
                                                                                             Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               core
                                                                                                                                                                                                                                                                                                                                   containing T-cell
against HIV-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein;
                                                           0,
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                                                           Gaps
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Human immunodeficiency virus

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RESULT 187
AAY39756
ID AAY397
XX AAY397
AC AAY397
XT 17-OCT
DT 17-OCT
DT 26-NOV
XX HIV1 c
XX HIV1 c
XX HIV5 V
KW infect
XX Human
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Best Local S
Matches 7
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                                                                                 17-OCT-2003
26-NOV-1999
                                                                                                                                                                                                                                                                                                                         The present invention describes a synthetic peptide comprising an amino acid sequence containing a T-cell epitope of an HIV gag protein linked a its C terminus to an amino acid sequence containing a B-cell epitope of an HIV gp41 protein and containing the amino acid sequence: XILKWMX2; where X1 = E, A, G or Q, and X2 = A or T, or an amino acid sequence capable of eliciting an HIV-specific antiserum and recognizing the sequence XILKDMX2. The synthetic peptide is useful in vaccines against HIV infection and in diagnostic applications. AAW99892 to AAW98966, and AAW99999 to AAW99989 represent synthetic peptides from the present
                                                                                                                  AAY39756;
                                                                                                                                          AAY39756
                                                                                                                                                                                                                                                                                             Sequence 17
                       infection; antibody; antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic chimeric HIV polypeptides - comprising epitope linked to gp41 B-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-189590/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Col 41-42; 41pp; English:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chong P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-JUN-1993;
09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV-1 vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW99958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW99958 standard; peptide; 17 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gag protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CONN-) CONNAUGHT LAB LTD.
                                     vaccine;
                                                          chimeric
                                                                                                                                                                                                   11
                                                                                                                                                                                                                                              l Similarity
7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human
                                                                                                                                         standard; peptide;
                                                                                                                                                                                                                          RAFVTIG 7
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                                                                                                                                                                                                                                                                                             A,
                                                                                                                                                                                                                                               Conservative
                                                                               (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunodeficiency virus; vaccine; T-cell
B-cell epitope; gp41 protein; chimeric;
                                  immunogenic composition;
                                                          peptide V3-LAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    synthetic peptide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93US-00073378
94US-00257528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95US-00462507
                                                                                                                                                                                                                                                                     87.2%;
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                                                                                                                                          17
                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                          Score 34;
Pred. No.
                                                                                                                                         A
                                                                                                                                                                                                                                              red. No. 2;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ID NO:35
                                  T cell
                                                                                                                                                                                                                                                                     'n
                                  epitope;
                                                                                                                                                                                                                                                                    Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  epitope;
infection.
                                  ₩
                                 cell epitope;
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RESULT 188
ADR18895
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Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                three-dimensional atomic structural conformation; protein co-ordinate data; V3 loop peptide; HIV-1; envelope glycoprotein; gp120; human monoclonal antibody 447-520; murine monoclonal antibody 0.5 beta; immunogen; immunogenic; with loop eptope; HIV-1 infectivity inhibitor; anti-HIV; vaccine;
                                                                                                                                                                                                                                                                       HIV-1 V3 loop related peptide SEQ ID NO:28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T cell epitopes and a sequence encoding 1 or more B cell epitopes and a carrier. Both the T cell and B cell epitopes are derived from HIV proteins. The compositions are useful as vaccines against HIV infection. The composition induces HIV-1-specific polyclonal antibodies that are opsonising and antiviral. The peptide components may be selected to induce a response against different viral isolates and in subjects who recognise different T cell epitopes. (Updated on 17-OCT-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence represents a fragment of a HIV1 protein, and can be used in the immunogenic composition of the invention. The composition comprises a synthetic fusion polypeptide which includes a sequence encoding 1 or more
                                                                                                                                                                                                                                                                                                     04-NOV-2004
                                                                                                                                                                                                                                                                                                                                                             ADR18895 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-550482/46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-JUN-1993;
09-JUN-1994;
               04-FEB-2003; 2003US-0444682P
                                        04-FEB-2004; 2004WO-US003304.
                                                                                                                              Synthetic.
                                                                                                                                            Human immunodeficiency virus l
                                                                                                                                                                        HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Col 22; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virus, useful antigens in producing vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Klein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-SEP-1999
                                                                                                 WO2004069863-A2
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                                                                                                                                                                        infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                     11 RAFVTIG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RAFVTIG 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                     (first entry)
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94US-00257528.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          87.2%; Score 34;
100.0%; Pred. No.
tive 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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CC isolated peptide molecule or isostere that minics the three-dimensional CC (3D) atomic structural conformation of the V3 loop peptide of the HIV-1 CC envelope glycopyretin gpl30 that is bound to, and constrained by, human CC differs in conformation from the same V3 loop peptide when it is in free CC differs in conformation from the same V3 loop peptide when it is in free CC differs in conformation from the same V3 loop peptide when it is in free CC compounds a molecule that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (3) a composition (C2) CC that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (3) a composition (C2) CC that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (3) a composition (C2) CC that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of thinding of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (3) a composition (C2) CC for induction of an anti-HIV-1 antibody response specific for a V3 loop eptope, comprising (C1) and an excipient; (5) a plantameceutical composition (C3) useful for blocking the interaction of HIV-1 with an R5 CC composition (C4) useful for blocking the interaction of HIV-1 with an R5 CC composition (C4) useful for blocking the interaction of HIV-1 with an R5 CC composition of an anti-HIV-1 V3 view peptide, (7) a computer generated composition of the peptide that is bound to 447-52D or 0.5beta Mbo or its antigen binding fragments, comprising a D acomic structure defined by NC; and composition (C2) is useful for inducing in a retrievable format. (C2) is useful for inducing in a subject an anti-HIV-1 conformation (C2) is useful for conformation of the 
Composition for inhibiting HIV-1 infection, comprises isolated peptide molecule that mimics atomic structural conformation of V3 loop peptide HIV-1 envelope glycoprotein that is bound to, and constrained by human monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a composition (C1) which comprises an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (YEDA ) YEDA RES & DEV CO LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8; SEQ ID NO 28; 127pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sharon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Schapira M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zolla-Pazner S,
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AAR38526
ID AAR3
XX
AC AAR3
XX
DT 25-N
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                                                  RESULT 189
                                                                                                             Matches
                                                                                                                    Query Match
Best Local (
25-MAR-2003
                 AAR38526;
                                 AAR38526 standard; peptide; 18
                                                                           11 RAFVTIG 17
                                                                                       1 RAFVIIG
                                                                                                            Similarity 7; Conserv
                                                                                                             Conservative
 (revised)
                                                                                            7
                                                                                                                    87.2%; Score 34; 100.0%; Pred. No.
                                                                                                             0
                                                                                                             Mismatches
                                                                                                                     2;
                                                                                                             0
                                                                                                                             Length 17
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0

Sequence 17

A.

exemplification

of the

present invention.

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RESULT 190
AAW03404
ID AAW034
XX
AC AAW034
XX
DT 10-OCT
XX
DE HIV px
XX
KW conjug
KW outer
                                                                                                                                                                     밁
                                                                                                                                                                                           S
                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 7
                                                                                                                                                                                                                                                                                     The sequences given in AAR38520-26 are cyclic HIV principle neutralising determinant (PND) peptides. They are stable compounds which, when conjugated with an immunological carrier, may be used to raise antipeptide, anti-HIV and HIV-neutralising antibodies. These may be used in vaccines for prevention or treatment of HIV-related diseases. They may also be used as reagents in the study of structure/function relationships in induction of HIV-neutralising immune response. (Updated on 25-MAR-2003 to correct PN field.)
  conjugate; PND;
outer membrane ]
                                    HIV principal neutralizing determinant cPND535
                                                                                    AAW03404;
                                                                                                                                                                                                                                                                                                                                                                                                                                Cyclic HIV principal neutralising determinant (PND) peptide(s) - useful as vaccines for treating and preventing HIV-mediated diseases e.g. AIDS
                                                           10-OCT-1996
                                                                                                          AAW03404 standard; peptide;
                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 26; 27pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bednarek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1993-228557/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-JAN-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EP551689-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cyclic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vaccine; immune response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-DEC-1993
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                                                                                                                                                                                           1 RAFVTIG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV PND
                                                                                                                                                                                                                  Similarity 7; Conserv
                                                                                                                                                                    RAFVTIG 18
                                                                                                                                                                                                                                                                  18
                                                                                                                                                                                                                                                                 A,
                                                                                                                                                                                                                  Conservative
                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      principle neutralising carrier; antipeptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tolman RL;
; HIV; principal r
protein complex;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptide cPND535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91US-00807943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92EP-00300243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     form cyclic peptide"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'note= "C-terminal condensed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note= "Abu =
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      label= OTHER
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                                                                                                                                                                                                                              87.2%;
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                                                                                                           18
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                                                                                                                                                                                                                              Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino group forms a cyclic peptide by
the C-terminal of Gly18"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gamma aminobutyric acid"
neutralizing determinant; OMPC;
; anionic spacer; vaccine;
                                                                                                           8
                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     determinant; PND; conjugate;
anti-HIV; HIV-neutralising; antibody;
                                                                                                                                                                                                                            DB 2;
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RESULT 191
ADR18878
ID ADR188
XX ADR188
AC ADR188
XX AT 04-NOV
XX HIV-1
XX Three-
XW Prote:
KW Prote:
KW Sp120,
KW W Slow
KW W HIV-1
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                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local
three-dimensional atomic structural conformation; protein co-ordinate data; V3 loop peptide; HIV-1; envelope g gp120; human monoclonal antibody 447-52D; murine monoclonal antibody 0.5 beta; immunogen; immunogenic; V3 loop epilope; HIV-1 infectivity inhibitor; anti-HIV; vacc HIV-1 infection.
                                                                                        HIV-1 V3-IIIB
                                                                                                                                                                                                                                                                                                                                                           A new conjugate immunogen comprises (a) the OMPC of Neisseria meningitidis b as a protein carrier, (b) a principal neutralideterminant (PND) of HIV as a peptidyl epitope against which responses are desired, and (c) a low mol. wt. anionic spacer and (b). The conjugate is water-soluble, yet can carry a high epitope loading. It is useful as a vaccine against HIV. The p sequence is an example of a PND used in the conjugate
                                                                                                                  04-NOV-2004
                                                                                                                                           ADR18878
                                                                                                                                                                   ADR18878 standard;
                                                                                                                                                                                                                                                                                                                                    Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 10; Page 70; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New conjugates of outer membrane protein and HIV epitope - for generat: HIV-neutralising response, have components joined by anionic spacer to ensure solubility of prod.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tolman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-MAY-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1994-128412/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human immunodeficiency virus; water-soluble
                                                                                                                                                                                                                                 12 RAFVTIG
                                                                                                                                                                                                                                                        1 RAFVTIG 7
                                                                                                                                                                                                                                                                                 l Similarity
7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RL,
                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Marburg
                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "one amino group of this lysine residue is linked to Neisseria meningitidis OMPC via a specified anionic spacer group; and the other forms a peptide bond with the terminal carboxy group of Gly(18), giving a cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "the
Lys(1). See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide"
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                                                                                                                                                                   peptide; 18
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                                                                                                                                                                                                                                                                                                            87.2%;
                                                                                        SEQ
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above"
                                                                                                                                                                                                                                                                                 Score 34; DB; Pred. No. 2.1
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                                                                                         NO:4.
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                                                                                                                                                                                                                                                                                                                                                                                                                          (a) the OMPC of Neisseria(b) a principal neutralizing
                                                                                                                                                                                                                                                                                                 2 B
                                                                                                                                                                                                                                                                                           .1;
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                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                    anionic spacer linking can carry a high peptide
                                                                                                                                                                                                                                                                                                         Length 18
                                                 envelope glycoprotein;
              vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       onto
                                                                                                                                                                                                                                                                                                                                                                        The present
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cc isolated peptide molecule or isostere that mimics the three-dimensional cenvelope glycoprotein gpl20 that is bound to, and constrained by, human collishment of the Mah, where the constrained V3 loop peptide of the HIV-1 collishment of the Mah, where the constrained V3 loop peptide of the HIV-1 collishment of the Mah, where the constrained V3 loop peptide when it is in free cities in conformation from the same V3 loop peptide when it is in free compounds a molecule that is useful as an HIV-1 V3 loop peptide when it is in free compounds a molecule that is useful as an HIV-1 V3 loop immunogen or as an inhibitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (2) designing compounds a molecule that is useful as an HIV-1 V3 loop immunogen or as an chimbitor of binding of HIV-1 to a chemokine receptor/HIV-1 co-receptor of that is useful as an HIV-1 V3 loop immunogen or as an chimbitor of chinding of HIV-1 to a chemokine receptor/HIV-1 co-receptor of that is useful as an HIV-1 V3 loop immunogen or as an chimbitor of composition (C3) useful for blocking the interaction of HIV-1 to a chemokine receptor/HIV-1 co-receptor on the surface of a receptor-bearing target cell; (3) a composition (C3) composition (C4) useful for blocking the interaction of HIV-1 with an R5 cc composition (C4) useful for blocking the interaction of HIV-1 with an R5 cc composition (C4) useful for blocking the interaction of HIV-1 with an R5 cc composition (C4) and an excipient; (6) a computer generated cc downward of the translational place peptide; (7) a computer generated cc downward translated and that includes a set of structure coordinates defining a B structure of a V3 loop peptide that is bound to 447-52D or 0.5beta Mab or its antigen binding fragment. (C1) has anticated and that includes a set of structure coordinates defining a B structure of a V3 loop peptide that is conformationally constrained by being bound cc 447-52D or 0.5beta Mab or its antigen binding fragment. (C1)
Matches
                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus Synthetic.
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                                                                                                                                                Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 4; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      monoclonal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2004; 2004WO-US003304
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   Local Similarity
les 7; Conserv
87.2%; Score 34; DB 8; ilarity 100.0%; Pred. No. 2.1; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             composition
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                                                                          Length 18;
   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ω
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   ٥,
   Gaps
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RESULT 192
AAP90279
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AAR25471
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X B X B B B X B X B X B
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                                                                                                                                                           Matches
                                                                                                                                                                            Query Match
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24-OCT-2003
25-MAR-2003
       V3 loop structure
                          25-MAR-2003
15-JAN-1993
                                                       AAR25471;
                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                  Revised record issued on 09-SEP-2004 : Correction to location
                                                                                                                                                                                                                                  Protein derivative stimulates a lymphocyte proliferative response in HIV-infected humans, providing a means of diagnosis, protection and therapeutic value. (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                          Claim 1; Page 27; 29pp; English
                                                                                                                                                                                                                                                                                                           New HIV proteins and peptide(s) therapy of AIDS, esp. for prepn
                                                                                                                                                                                                                                                                                                                                                          Rusche JR,
Petro J, (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Simian-Human
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                                                                        AAR25471
                                                                                                                                                                                                                                                                                                                                        WPI; 1989-070387/10.
                                                                                                                                                                                                                                                                                                                                                                                                         27-AUG-1987;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified
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                                                                                                                                                                    Local
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                                                                                                                                                         Similarity 7; Conserv
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                                                                                                                                     RAFVTIGK 8
                                                                                                                                                                                               20
                                                                       standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; protein; 20
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                                                                                                                       RAFFTIGK 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             env gene; HIV vaccine; ds
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keeffe T;
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                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunodeficiency virus.
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(revised)
(revised)
                          (revised)
(first entry)
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                                                                                                                                                                   87.2%;
                                                                                                                                                                                                                                                                                                                                                                    Jayaherian K,
                                                                                                                                                                                                                                                                                                            prepn.
                                                                          20
                                                                                                                                                           Score 34; DB Pred. No. 2.3; 0; Mismatches
                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                           used in diagnosis,
of vaccines against
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                                                                                                                                                                    DB 1;
2.3;
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                                                                                                                                                                            Length 20
                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                              prophylaxis or HIV infection.
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RESULT 194
AAW76842
ID AAW768
AC AAW
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                 B cell; T cell; epitope; immunoglobulin; heavy chain; gpl20; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The peptide sequence given represents the sequence from amino acid 310-328 of the external protein gp120 from HIV. This comprises a disulphide loop in the third variable region. It was used in an example of the invention and was incorporated into hepatitis B surface antigen (HBsAg) particles. The hybrid formed in this reaction is useful as a vaccine for the prophylactic treament of various infectious diseases eg. AIDS. Conjugation of this peptide with the HBsAg particle allows its processing to be directed via a non-endosomal route. In this way the gp120 fragment can be recognized by cytotoxic lymphocytes. (Updated on 25-MAR-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New immunogenic hybrid polypeptide(s) for vaccine formulations - cor S antigen of hepatitis B linked via spacer to heterologous antigen, gD from HSV or gp120 form HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis B; surface antigen; AIDS; cytotoxic lymphocytes; disulphide loop; variable region.
                               13-FEB-1998;
                                                                                   20-AUG-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Fusion immunoglobulin heavy chain HIV gp120 B cell epitope #12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW76842 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 AA;
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21-MAR-1991;
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                                                                                                                                   WO9836087-A1
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                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                      gp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RAFVTIG 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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91GB-00005993.
                               98WO-US002766
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                87.2%; but
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 34; DB 2; Pred. No. 2.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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en, e.g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                               02-JUL-2001; 2001GB-00016151
29-NOV-2001; 2001GB-00028638
28-JAN-2002; 2002GB-00001896
28-MAR-2002; 2002GB-00007509
                                                                                                                                            02-JUL-2002; 2002WO-GB003037.
                                                                                                                                                                                                                                           Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                          proliferative immune response; antiinflammatory; neuroprotective;
cytostatic; gene therapy; vaccine; inflammatory disease; nerve damage;
autoimmune disease; axonal damage; cancer; inflammatory; HIV.
                                                                                                                                                                                                                                                                                                                                                        HIV gp120 V3 loop derived peptide ARP740/28.
                                                                                                                                                                                                                                                                                                                                                                                          23-OCT-2003
14-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                        ABP57070;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABP57070 standard; peptide; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells, used to induce tolerance to the epitopes for treatment of human immune deficiency
 Dalgleish AG,
                                                                                                                                                                            16-JAN-2003.
                                                                                                                                                                                                             WO2003004049-A2
                                                                                                                                                                                                                                                                                                                          Anti-human leukocyte antigen antibody; anti-HLA antibody; anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 10; Page 119; 154pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-506315/43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Scott D,
                                 (ICEB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14 RAFVTIG 20
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                                 ICE BIOLOGICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first en
 Cadogan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-0040581P.
                                                                                                                                                                                                                                                                                                                                                                                        entry)
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                                 LTD.
 Heeney J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
 White SDT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
2.3;
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WPI; 2003-210314/20

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RESULT 196
AAR66425
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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an anti-human leukocyte antigen (HLA) antibody (I) useful for the preparation of a medicament for treating a disease involving a proliferative immune response. (I) has anti-HIV, antiinflammatory, neuroprotective and cytostatic activities, and can be used in vaccines and in gene therapy. The antibody (I) is useful for the preparation of a medicament for treating diseases involving a proliferative immune response, e.g. HIV, inflammatory diseases, autoimmune diseases, axonal or nerve damage or related impairment or cancers, and other diseases or conditions with an inflammatory component. The present sequence represents an HIV gp120 V3 loop derived peptide, which is used in the exemplification of the present invention. (Updated on 23-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                             T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Use of anti-HIA antibody for the preparation of a medicament for treating a disease involving a proliferative immune response e.g. HIV, inflammatory diseases, autoimmune diseases, axonal/nerve damage or
                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                      HIV-1 IIIB
                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
03-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR66425;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 AA;
                                                                            Polypeptide inducing helper T cell, cyt
responses - to target antigen in hosts
                                                                                                                   WPI; 1995-006707/01.
                                                                                                                                                                                               14-MAY-1993;
                                                                                                                                                                                                                        13-MAY-1994;
                                                                                                                                                                                                                                                    24-NOV-1994.
                                                                                                                                                                                                                                                                             WO9426785-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR66425 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 41; 69pp; English.
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                                                               for therapeutic
                                                                                                                                                                     (USSH ) US SEC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     impairment, cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RAFVTIG
                                                                                                                                          JA,
                                                                                                                                                                                                                                                                                                                                                                                   peptide 18-10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
(first entry)
                                                              target antigen in hosts of
c or prophylactic vaccines
                                                                                                                                          Ahlers
                                                                                                                                                                      DEPT HEALTH.
                                                                                                                                                                                               9308-00060988
                                                                                                                                                                                                                         94WO-US005142
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87.2%; but
100.0%; Pr
                                                                                                                                          'n
                                                                                                                                          Pendleton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                          cytotoxic T cell and antibodies sts of different MHC haplotypes,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 6
                                                             against
                                                                                                                                            Nara
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                                                                             egp.
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Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residu

residue

Sequence 15 AA;

Example 1; Page

33;

120pp; English

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RESULT 197
AAR66429
ID AAR664
XX AAR664
XX AAR664
XX DT 25-MAR
DT 25-MAR
DT 03-AUG
XX T cell
XX T cell
XW Cluste
XX Cluste
XX Synthe
XX W09426
XX 14-MAY
XX 14-MAY
XX T respon
PT respon
PT for th
XX WPI; 1
XX WPI; 1
XX WPI; 1
XX WPI; 1
XX WFI; 1
XX Cluste
X
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                                                                                                                                                                                                                       Single residues from the HIV-1 RF sequence (AAR66415) replaced residues in the HIV-1 IIIB sequence (AAR66414) to test the effect of each residue on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see AAR66409 and AAR66411). Analysis of the substituted peptides (AAR66416-R66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups of the substraction of the substractio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       on the binding of neutralising and non-neutralising sera from animals immunised with the cluster peptides PCLUS 3-18 and PCLUS 6-18 (see ARR66409 and ARR66411). Analysis of the substituted peptides (ARR66416-R66430) showed that binding was enhanced over peptide 18 control when a tyrosine was substd. for a Val at position 11 and substns. at positions 12,13,14 and 15 revealed a similar enhancement. Binding of both groups o sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-10, the Phe residue at position 10 in peptide 18 has been replaced by a Ile residue. (Updated o 25-MAR-2003 to correct PN field.)
                                                                           sera was reduced when substns. were made in the principal neutralising determinant sequence (PGRAF). In peptide 18-14, the Gly residue at position 14 in peptide 18 has been replaced by an Ala residue. (Update on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T cell helper site; cytotoxic T cell response; neutralising antibody; human immunodeficiency virus type 1; envelope glycoprotein gp120; cluster peptide; principal neutralising determinant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polypeptide inducing helper T cell, cytotoxic T cell and responses - to target antigen in hosts of different MHC report therapeutic or prophylactic vaccines against HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1995-006707/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Berzofsky JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV-1 IIIB peptide 18-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003
03-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 33; 120pp; English.
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(first en
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Length 15

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RESULT 199
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ID AAO156
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ABB05778
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Best Local S
Matches 7
                                                                                                                                              Matches
                                                                                                                                                         Query Match
Best Local (
                                                                                                                                                                                                                 are: (1) a method for producing N1 and artificial gene expression vectors comprising N1; (2) transgenic non-human animals comprising N1; and (3) treatments and diagnostic reagents containing an artificial protein, artificial tissues or high molecular weight artificial proteins. N1 is useful for creating industrially useful artificial artix proteins for cell culture. The present sequence represents a peptide which is used in an example from the present invention. (Updated on 29-AUG-2003 to standardise OS field)
AA015660;
                       AAO15660 standard; peptide; 7 AA
                                                                                                                                                                                               Sequence 7
                                                                                                                                                                                                                                                                                                                             The present invention describes a polyfunctional base sequence having two or more functions in different reading frames. Also
                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 47; 61pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                      Polyfunctional base sequence having two or more functions in different reading frames, useful for producing artificial matrix proteins for cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JUN-2001; 2001WO-JP005116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-AUG-2003
07-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-098069/13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200196558-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polyfunctional base sequence; microgene; industrial; cell culture; artificial matrix protein; transgenic animal; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV gp120 related peptide SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB05778;
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                                                                                                                                            Similarity 6; Conserv
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                                                                                            FVTICK 6
                                                                                                                  FVTIGK 8
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                                                                                                                                                                                                 B
                                                                                                                                            Conservative
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(first entry)
                                                                                                                                         76.9%; Score 30; DB 5; L
100.0%; Pred. No. 1.8e+06;
tive 0; Mismatches 0;
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Pred. No. 2.9;
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                                                                                                                                                                  Length 7;
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RESULT 200
AAW38251
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AX 19-MAR
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KW Omega
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Best Local Similarity
                                                                                                                                      Multivalent chimeric peptide; tandem repeat unit; human; mucin 1; Nomega loop sequence; prophylaxis; therapy; GPGR crest; pathogenic virus neutralisation; human immunodeficiency virus; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention comprises an artificial protein which induces a strong immune response to a peptide group (the protein contains all or part of the peptide group). The artificial protein assists the formation of high-order protein structure and/or assists the antigen presentation of immunocompetent cells. The artificial protein of the invention is useful for inducing a strong immune response and the preparation of effective antibodies to specific antigens, especially HIV. The present amino acid sequence represents a peptide that was used in the invention
31-JAN-1996;
                             29-JAN-1997;
                                                                                                                                                                                                                             19-MAR-1998
                                                         07-AUG-1997.
                                                                                                              Homo sapiens
                                                                                   WO9728187-A2
                                                                                                                                                                                               Carboxy-terminal of GPGR crest.
                                                                                                                                                                                                                                                         AAW38251;
                                                                                                                                                                                                                                                                                   AAW38251 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 46; 77pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   other antigens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Artificial protein capable of inducing a strong immune response to peptide group for assisting antibody production in vivo to viruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-519151/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shiba K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-OCT-2000; 2000JP-00314288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-OCT-2001; 2001WO-JP008893
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NISC-) JAPAN SCI & TECHNOLOGY CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-APR-2002.
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                                                                                                                                                                                                                                                                                                                                                                                   ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immune response induction; high-order protein structure formation;
                                                                                                                                                                                                                                                                                                                                                                                                            6;
                                                                                                                                                                                                                                                                                                                                                                        FVTIGK 8
                                                                                                                                                                                                                                                                                                                                                       FVTIGK 6
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                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                           (first entry)
                             97WO-US001726.
                                                                                                                                                                                                                                                                                  peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                         76.9%; Score 30; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 7;
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                                                                                                                                                                   1; MUC1;
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15-OCT-1996;

96US-00730244.

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RESULT 201
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Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B cell; T cell; epitope; immunoglobulin; heavy chain; gp120; IgH; human immune deficiency virus; HIV; tolerance; treatment; therapy; prophylaxis; vaccine; chemotherapy; immune response; modifier; tumour; microbial infection; autoimmune disease; antibody; apoptosis; antiviral T cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New multivalent chimeric peptide(s) for neutralising pathogenic microbes - comprising a loop structure of human mucin 1 and an omega loop of an immunoglobulin superfamily protein.
New fusion immunoglobulin heavy chain including gp120 epitopes and related complete antibodies - DNA, vectors and transformed cells,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fusion immunoglobulin heavy chain HIV gp120 T cell epitope #40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW76971 standard; peptide; 15
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                                                                                    WPI; 1998-506315/43
                                                                                                                                                                                                                                                        13-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                    20-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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ص
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                                                                                                                                         Zambidis
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100.0%; Pred. No.
tive 0; Mismatc
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RESULT 202
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XX pri
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
"---hes 6; Conservative
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                                                          Cunningham B,
                                                                                                                                                                                                                                         07-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                           GB2282813-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             annular antigen scaffold core; AASC; HIV V3 loop; lysine; principal neutralising determinant; PND; cyclic; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cyclic HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR90229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR90229 standard; peptide; 15 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       induce tolerance to the epitopes for treatment of human immune deficiency virus infection.
                                                                                                                                                                               15-OCT-1993;
                                                                                                                                                                                                                                                                                                 19-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified-site
                                                                                                                       (MERI ) MERCK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 FVTIGK 8
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                                                                                                                       S
                                                             Hannah J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   peptide attached to annular antigen scaffold
                                                                                                                                                                                  93US-00138514
                                                                                                                                                                                                                                            94GB-00020263
                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "this is an epsilon-Lys residue, the alpha-amino and carboxy terminals of which are incorporated into an annular antigen scaffold core of formula KKKKCX as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "This residue is bonded to the thiol sulphur of Cys(13) via a -CO-CH2- linkage, formed by introducing a bromeacetyl group onto the N-terminal and allowing the Br to condense with the Cys side chain"
                                                                                                                                                                                                                                                                                                                                                                                                                      described in AAR90224"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "see above"
15
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                                                                Tolman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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Pred. No.
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AAW32887
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New annular antigen scaffold cores are provided for antigens or epitopes such as HIV V3 loop peptides (e.g. the present sequence; but see also GB2282815; AAR90219 - AAR90223), GRRH peptides, malaria antigenic peptides or bacterial capsular polysaccharides. The scaffolds comprise a ring of 3-10 Lys residues cyclised via a thioether linkage. The epitopes or antigens are bonded to each of the Lys side-chain amino groups. The Cterminus of the scaffold may be linked to a moiety such as beta-alanine or a peptide providing a T cell epitope, a lipopeptide which may provide an adjuvant effect, or another molety providing a carrier function. The scaffolds constitute effective synthetic vaccines. The present sequence represents one of four identical thioether-cyclised HIV V3 loop peptides which are attached to each of the four Lys residues in the the annular
                                                                                                 Synthetic human immunodeficiency virus vaccine - comprising hydrophilic peptide corresponding to at least 1 antigenic determinant of envelope
                                                                                                                                                WPI; 1997-244862/22
                                                                                                                                                                                                                               20-OCT-1995;
09-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                          Hydrophilic; antigenic determinant; HIV; envelope; glycoprotein; env; gp, recognition; B lymphocyte; type specific; antibody; vaccine; protection;
                                                                                                                                                                                                                                                                        18-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-JAN-1998
                                                                                                                                                                                                     (-uaxn)
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                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV envelope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW32887 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            scaffold core described in AAR90224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New poly:lysine annular core for carrying epitope(s) peptide, gonadotropin releasing hormone, malarial or useful in vaccines.
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                                                                                                                                                                                                       VIND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RAFVTIGK
                                                                                                                                                                           Palker TJ;
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llarity 75.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
                                                                                                                                                                                                                                                                                                                                                                                                                                     glycoprotein 120 T cell epitope P10.
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96US-00599266.
                                                                                                                                                                                                                                                                        96WO-US016911.
                                                                                                                                                                                                                                                                                                                                                                                infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                                                                                                                                                neutralisation; epitope.
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Pred. No. 23;
0; Mismatches
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    esp HIV
    bacterial

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peptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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An essentially pure hydrophilic peptide, comprising at least 1 antigenic determinant of human immunodeficiency virus (HIV) envelope (env) glycoprotein (gp) recognised by B lymphocytes, when covalently linked to a carrier molecule, i.e. the present sequence, induces the production of

S

RAFVTIG

Query Match Best Local S Matches 5

Similarity 5; Conserv

Conservative

74.4%;

Score 29; DB Pred. No. 23; 1; Mismatches

۳.

0;

Gaps

0

BB

2

Length 15;

peptide corresponding to at le glyco:protein recognised by B

Disclosure;

Page

27;

104pp; English.

lymphocytes

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RESULT 204
AAW55971
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Best Local
                   The map kinase kinase 3 (MKK3) N-terminal peptide was used to raise polyclonal anti-MKK3 antibodies. These antibodies were used in the immunoprecipitation of stress-activated protesin kinases (SAPK). The invention claims for the human SAPK4 cDNA (AAV26081) isolated from a human pituitary gland cDNA library. The invention also claims that SAPK4 protesin (AAW35967) can be useful in a screening assay for identifying agents that inhibit its activity and/or agents that block its activation through stress activated kinase kinase 3 (SKK3). Therefore, the agents identified in the assays may be potentially useful for treating inflammatory diseases, e.g. rheumatoid arthritis, asthma and psoriasis
Sequence 15
                                                                                                                                                                                                             New stress-activated protein kinase 4 - e.g. anti-inflammatory, anti-cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stress-activated protein kinase 4; SAPK4; pituitary gland; asthma; stress activated kinase kinase 3; SKK3; rheumatoid arthritis; psor: inflammatory disease; immunoprecipitation; MKK3; map kinase kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       high titres of protective, type specific anti-HIV antibodies (Ab) in a mammal. The peptide can be used in vaccines for producing a protective immune response to HIV infection, while a HIV neutralising Ab can be induced in a primate by administering a composition comprising HIV env peptides that disrupt gp120/gp41 interactions
                                                                                                                                                                                                                                                          WPI; 1998-240806/21
                                                                                                                                                                                                                                                                                     Cohen
                                                                                                                                                                                                                                                                                                                                             09-OCT-1996;
15-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                       09-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                   16-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9815618-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Map kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW55971;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15
                                                                                                                                                                                                                                                                                                                (MEDI-) MEDICAL RES COUNCIL.
                                                                                                                                                                                                                                                                                     P,
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                                                                                                                                                                                  4; Page 63; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
7; Conserva
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                                                                                                                                                                                                                                                                                     Goedert M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           kinase 3
ΑA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                             96GB-00021096.
97GB-00009781.
                                                                                                                                                                                                                                                                                                                                                                                       97WO-GB002779.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MKK3) N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         74.4%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide
                                                                                                                                                                                                               useful in drug screening, immuno-suppressing agents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             psoriasis;
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RESULT 206
AAW25898
ID AAW258
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AC AAW258
AC AAW258
AC TO AAW258
XX
DT 25-MAR
DT 22-OCT
XX
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AAR68680
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                                                                                                                                                                                               Matches
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Best Local (
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25-MAR-2003
07-SEP-1995
                                                                                                                                                                                                                                                                                    This sequence represents a B-cell epitope sequence derived from the V3 loop of the HIV-1 isolate, LAI. This B-cell epitope may be linked to a T-cell epitope also derived from HIV-1. These chimeric peptides may then be used in the production of HIV-1 vaccines. These peptides in which the peptides are C-terminally modified by the addition of a Lys residue which is modified on its epsilon amino acid to carry an additional copy of the peptide molecule. The linear and multimeric peptides may be used for the treatment of AIDS by acting to displace the binding of HIV virus to human or animal cells or by disturbing the 3D organisation of the virus.

(Updated on 25-MAR-2003 to correct PN field.) (Updated on 16-CCT-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel tandem synthetic HIV-1 peptide(s) - comprising T-cell epitope gag protein linked to B-cell epitope of V3 loop protein of an HIV-I isolate.
               25-MAR-2003
22-OCT-1997
                                                                                                                                                                                                                                                  Sequence 20 AA;
                                                                                                                                                                                                                                                                            (Updated on 
standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 16; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T-cell; epitope; HIV-1; core protein; p24E; B-cell; antigen; pol; vaccine; multimeric peptide; AIDS; 3D organisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  B cell epitope, LAI.
                                                    AAW25898;
                                                                              AAW25898 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sia CDY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-JUN-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR68680 standard; peptide; 20 AA
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                                                                                                                                              14
                                                                                                                                                                     1 RAFVTIG 7
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                                                                                                                                                                                              Similarity 6; Conservat
                                                                                                                                              RAFYTIG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chong P,
                                                                                                                                                                                                                                                                            25-MAR-2003
OS field)
                                                                                                                                                                                                 Conservative
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(revised)
(first entry)
            (revised)
(first entry)
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                                                                                                                                             20
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                                                                              peptide; 20 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Klein MH
                                                                                                                                                                                                            74.4%;
85.7%;
                                                                                                                                                                                               <u>.</u>
                                                                                                                                                                                                            Score 29; DB
Pred. No. 31;
                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                         DB 2;
                                                                                                                                                                                               1; Indels
                                                                                                                                                                                                                        Length 20;
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                                                                                                                                                                                              Gaps
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RESULT 207
AAW25850
ID AAW258
XX AAW258
XX AAW258
XX 25-MAR
DT 20-OCT
XX HIV: h
XX HIV; h
KW V3 loc
XX V3 loc
XX Synthe
XX V3 loc
XX Synthe
XX U5563:
XX O9-JUh
XX O9-JUh
XX O9-JUh
XX O9-JUh
XX O9-JUh
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to new synthetic peptides comprising at least one amino acid sequence comprising an HIV gag protein T-cell epitope linked at its C- or N-terminus to an amino acid sequence comprising a B-cell epitope of the V3 loop of an HIV env protein, which can be used to generate vaccines against HIV-1. The T-cell epitope sequence is pref. selected from the T-helper determinant core peptides P24B, P24N, P24N, P24N and P24H while the B-cell epitopes are derived from HIV strains including CTLB-56, V3NN, CTLB-29, CTLB-55, SF2, LAI, IIIB, RF, Z6, 2054, 1714 and BX08. The peptides are chimaeric and can be linked to a branched Lys backbone. This sequence represents the HIV-1 strain IIIB env protein V3 loop B-cell epitope. (Updated on 25-MAR-2003 to correct PF field.)
                                          09-JUN-1994;
                                                                                                                               Synthetic.
                                                                                                                                                        HIV; human immunodeficiency virus; gag; T-cell; B-cell; epitope; env;
V3 loop; vaccine; determinant; chimaeric.
                                                                                                                                                                                                     HIV-1 strain LAI env protein V3 loop B-cell epitope.
                                                                                                                                                                                                                                  25-MAR-2003
20-OCT-1997
                                                                                                                                                                                                                                                                              AAW25850;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tandem synthetic HIV peptide(s) useful as immunogens - comprising gag protein T-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-332082/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV; human immunodeficiency virus; gag; T-cell; B-cell; epitope;
V3 loop; vaccine; determinant; chimaeric.
                                                                       17-JUN-1997
                                                                                                  US5639854-A
                                                                                                                                                                                                                                                                                                          AAW25850 standard; peptide; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 8; Col 74; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Klein MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-JUN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV-1 strain IIIB env protein V3 loop B-cell epitope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                 14
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6; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                            1 RAFVTIG 7
                                                                                                                                                                                                                                                                                                                                                                                 RAFYTIG 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                 (revised)
(first entry)
                                            94US-00257528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93US-00073378
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         74.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chong P;
                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 29;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB
31;
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09-JUN-1993;

93US-00073378

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RESULT 208
AAW67366
ID AAW673
XX AAW673
XX 17-OCT
DT 25-JAN
XX Immuno
XW Immuno
XW Immuno
XW V3 loc
XX V3 loc
XX V3 loc
XX U55817
XX 05-JUN
XX WPI 1
XX WPI 1
XX Synthe
PT Synthe
PT Synthe
PT Synthe
PT Synthe
CC Vaccir
CC vaccir
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            at its C- or N-terminus to an amino acid sequence comprising a B-cell epitope of the V3 loop of an HIV env protein, which can be used to generate vaccines against HIV-1. The T-cell epitope sequence is pref. selected from the T-helper determinant core peptides P24E, P24N, P24L, P24M and P24H while the B-cell epitopes are derived from HIV strains including CTLB-56, V3MM, CTLB-29, CTLB-55, SF2, LAI, IIIB, RF, Z6, Z054, 1714 and BX08. The peptides are chimaeric and can be linked to a branched Lys backbone. This sequence represents the HIV-1 strain LAI env protein V3 loop B-cell epitope. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                 Synthetic human immunodeficiency virus-1 peptide(s) - containing T-cell epitope and B-cell epitope(s) are candidate vaccines against HIV-1.
                                                                                                                                                                                                            09-JUN-1993;
09-JUN-1994;
                                                        Disclosure; Col
                                                                                                                                                                                                                                                    05-JUN-1995;
                                                                                                                                                                                                                                                                                 06-OCT-1998.
                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                        HIV-1 strain
                                                                                                                                                                                                                                                                                                                                                                                                                                  17-OCT-2003
25-JAN-1999
                                                                                                                            WPI; 1998-556461/47
                                                                                                                                                                                 (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                            US5817754-A
                                                                                                                                                                                                                                                                                                                                                                            Immunogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW67366;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW67366 standard; peptide; 20 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tandem synthetic HIV peptide(s) useful as immunogens -
protein T-cell epitope linked to env protein B-cell epi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to new synthetic peptides comprising at least one amino acid sequence comprising an HIV gag protein T-cell epitope linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Col
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RAFVTIG
                                                                                                                                                       Klein
                                                                                                                                                                                                                                                                                                                                                                         vaccine; HIV-1; T-cell; B-cell; epitope; core protein; gp120.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first ent
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94US-00257528.
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                                                     9; 40pp;
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                                                                                                                                                                                                                                                                                                                                                                                                     loop peptide epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
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                                                                                                                                                       Sia CDY
                                                       English
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הי
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Pred. No. 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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The invention relates to a novel immunogenic composition for use in vaccines for the treatment of HIV-1 comprising an HIV-1-derived T-co epitope linked to an HIV-1-derived B-cell epitope. The T-cell epitop

ed T-cell epitopes

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14

Query Match Best Local S Matches 6

6; Conserv

Conservative

0;

1;

Indels

<u>,</u>

Gaps

0

74.4%; 85.7%;

Score 29; DB Pred. No. 31; 0; Mismatches

DB 31;

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Length 20

RAFVTIG 7 RAFYTIG

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RESULT 209
AAM99974
ID AAM99974
XX AAM999
XX JAM999
CC JAM999
CC HIVent
CC JAM999
CC HIVent
CC JAM999
CC JAM99
CC JAM9
CC JAM99
CC JAM9
CC JAM99
CC JAM9
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Best Local Similarity
Thes 6; Conserve
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                                                                                                       The present invention describes a synthetic peptide comprising an amino acid sequence containing a T-cell epitope of an HIV gag protein linked a its C terminus to an amino acid sequence containing a B-cell epitope of an HIV gp41 protein and containing the amino acid sequence: XILKDMX2; where X1 = E, A, G or Q, and X2 = A or T, or an amino acid sequence capable of eliciting an HIV-specific antiserum and recognizing the sequence XILKDMX2. The synthetic peptide is useful in vaccines against HIV infection and in diagnostic applications. AAW98892 to AAW98906, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  are generally designed based on the p24 core protein and the B-cell epitopes from the V3 loop of the gp120 protein from various HIV-1 strains. This peptide corresponds to the V3 loop peptide epitope from HIV-1 strain LAI. The peptide is used to generate a hybrid T- and B-ce epitope (AAW67353). (Updated on 17-OCT-2003 to standardise OS field)
     Sequence 20
                                                                                 AAW99899 to
                                                                                                                                                                                                                                                                                                                                                     Example 1; Col 49-50;
                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic chiméric HIV polypeptides - comprising epitope linked to gp41 B-cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-189590/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-JUN-1993;
09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JUN-1995;
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gag protein;
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                                                          Invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Klein MH,
     AA;
                                                                              AAW99989 represent synthetic peptides from the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunodeficiency virus; vaccine; T-cell
B-cell epitope; gp41 protein; chimeric;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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94US-00257528.
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85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LTD.
                                                                                                                                                                                                                                                                                                                                                  41pp; English.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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infection.
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s field)
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AAR68800
ID AAR0
XX
AC AAR0
XX
DT 25-N
DT 23-J
XX
DE Cytc
XX
                                                                                                                                                                                                                                                                          RESULT 210
AAY39699
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                                                                                                                                                                                             8
                                                                                                                                RESULT 211
                                                                                                                                                                                                                     Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                           This sequence represents a fragment of a HIV1 protein, and can be used in the immunogenic composition of the invention. The composition comprises a synthetic fusion polypeptide which includes a sequence encoding 1 or more T cell epitopes and a sequence encoding 1 or more B cell epitopes and a sequence encoding 1 or more B cell epitopes and a carrier. Both the T cell and B cell epitopes are derived from HIV proteins. The compositions are useful as vaccines against HIV infection. The composition induces HIV-1-specific polyclonal antibodies that are opsomising and antiviral. The peptide components may be selected to induce a response against different viral isolates and in subjects who recognise different T cell epitopes. (Updated on 17-0CT-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Col 9; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    containing both the T and B cell epitopes of virus, useful antigens in producing vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-JUN-1993;
09-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5951986-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY39699;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY39699 standard; peptide;
          Cytotoxic T lymphocyte epitope 57 derived
                                      25-MAR-2003
23-AUG-1995
                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-550482/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Klein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-OCT-2003
26-NOV-1999
                                                                             AAR68800;
                                                                                                                                                                                                                                                                                                   standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunogenic composition containing synthetic fusion polypeptides containing both the T and B cell epitopes of the human immunodeficiency
                                                                                                    AAR68800 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CONN-) CONNAUGHT LAB LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ĭ,
                                                                                                                                                                      14
                                                                                                                                                                                                                       6; Conserv
                                                                                                                                                                      RAFYTIG
                                                                                                                                                                                          RAFVTIG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chong P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ne; immunogenic composition; T cell epitope; B cell epitope,
antibody; antiviral.
                                                                                                                                                                                                                                                                          ₽,
                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first entry)
                                      (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide LAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93US-00073378.
94US-00257528.
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                                                                                                                                                                       20
                                                                                                                                                                                                                                    74.4%;
85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                    Score 29; DB
Pred. No. 31;
                                                                                                      ₿
                                                                                                                                                                                                                         Mismatches
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             from
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                                                                                                                                                                                                                        1;
              env
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                                                                                                                                                                                                                         Indels
            gp120
             protein
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RESULT 212
AAY10165
ID AAY101
XX AAY101
XX AAY101
XX I2-MAY
DT 12-MAY
XX Cytoto
KW immuni
KW immuni
KW malign
XX Mo9902
XX WO9902
XX PD 21-JAN
XX PF 10-JUI
XX AAY10165
                                                                                                                                                                                                                                                                                                                                                                                            닭
                                                                                                                                                                                                                                                                                                                                                                                                                      S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR68744-805 are cytotoxic T lymphocyte (CTL) class I and II restricted epitopes derived from human immunodeficiency virus proteins. AAR68800 corresponds to amino acid residues 312-318 of the env gpl20 protein. These antigens are examples of peptides that can be used with an immunogenic implant. The implant is associated with an antigen associated with a pathogen and used to form a discrete, localised chronic inflammation site which acts as a local 'factory' for prodn. of CTL's which lyse cells infected with a specific pathogen. The expanded set of pathogen-specific CTL's can eradicate or prevent development of infection, and can also be used to treat or arrest the development of cancers associated with infection. (Updated on 25-MAR-2003 to correct PN cancers associated with infection.
                                                                                                                                                        Cytotoxic T-lymphocyte response; CTL; antigen; lymphatic system; immunisation; tumour; infectious disease; immunotherapy; cancer; malignant melanoma; viral disease; hepatitis; AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Implant for sustained release of pathogen-associated antigen - for chronic inflammatory site producing cytotoxic T-lymphocytes lysing infected cells, esp. for treating AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytotoxic T lymphocyte; epitope; antigen; pathogenic; nef; gag; pol; env; gp120; gp41; HIV; cell-mediated immunity; human immunodeficiency virus; class II restricted.
                                              21-JAN-1999.
                                                                                                                            Synthetic.
                                                                                                                                                                                                                     T cell epitope/MHC ligand SEQ ID NO:95.
                                                                                                                                                                                                                                                                                   AAY10165;
                                                                                                                                                                                                                                                                                                                AAY10165 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 12; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-036067/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leonard RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9428871-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus.
                                                                                                                                                                                                                                                    12-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ENDO-) ENDOCON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                          immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                        1 RAFVTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                            RAFVII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       71.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 28; DB; Pred. No. 1.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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10-JUL-1998;

98WO-US014289

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ARBSULT 213
ABG79847
ID ABG798
XX ABG798
XX ABG798
XX ABG798
XX ABG798
XX Major
C1
XX Major
KW Major
KW epitop
KW antige
KW 1ympho
KW acquir
XX AC
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a differentiation antigen, a tumour-specific multilineage antigen, an embryonic antigen, an oncogene antigen, a mutated tumour-suppressor gene antigen, or a viral antigen. They can be used for the treatment of disease such as cancer, e.g. malignant melanoma or infectious disease, e.g. viral disease such as hepatitis or AIDS. Sustained antigen delivery to the lymphatic system provides for potent CTL stimulation that takes place in the milieu of the lymphoid organ, and it sustains stimulation that is necessary to keep CTL active, cytotoxic and recirculating through the body. AAY10071 to AAY10639 represent examples of peptide antigens given in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal. The method comprises: (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal's lymphatic system to maintain the immunologic CTL response. The method can be used for the delivery of e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               epitope; cytotoxic T lymphocyte response; CTL response; lymphatic system; antigen; immunogenic; malignant tumour; carcinoma; melanoma; leukaemia; lymphoma; infectious disease; hepatitis; malaria; measles; tuberculosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inducing a cytotoxic T lymphocyte response - by maintaining a level of antigen in the lymphatic system of a mammal so as to provide a sustained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-JUL-1997;
10-DEC-1997;
                              Kundig
                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABG79847;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABG79847 standard; peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-120514/10
                                                                                                                                                                                                                            22-JAN-2002; 2002WO-US002033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Major histocompatibility complex; MHC; MHC class I molecule; virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MHC class I molecule, viral epitope #95.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-NOV-2002
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                                                                                              (CTLI-) CTL IMMUNOTHERAPIES CORP.
                                                                                                                                                                                                                                                                                                15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     response, used to treat, e.g. AIDS.
                                 ¥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 RAFVTI 6
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                                 Simard JJL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Simard JJL;
                                                                                                                                                               2001US-00776232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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97US-00988320.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     199pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2; bc.,
o. 1.8e+06;
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RESULT 214
ABR55441
ID ABR5544
XX ABR554
XX ABR554
XX ABR554
XX ANTige
XX ANTige
XX HUMAN
XX HUMAN
XX HUMAN
XX COMMAY
XX COMMA
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                                                                                                                                                                                            Novel composition for reducing interactions between Bob and gp120 and reducing symptoms of HIV enteropathy, HIV nephropathy or HIV-related infertility, comprises a Bob inhibitor that binds a region of Bob.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antigen; Bob; gp120; lymphocyte; HIV enteropathy; HIV nephropathy; HIV-related hyperlipidemia; HIV-related infertility.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR55441 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (a) an antigen in the form of a polypeptide; (b) a vector comprising nucleic acid encoding the antigen; or (c) a non-peptide antigen. The method is useful for inducing and/or sustaining CTL response in a man This is particularly useful for treating a mammmal having a malignant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal comprising administering directly to the lymphatic system of the mammal:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Inducing or sustaining immunological cytotoxic T lymphocyte response in ammmal, useful for treating a mammal with malignant tumor or infectious disease, by directly administering an antigen to the lymphatic system of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-657506/70
                                                                                                                                                                                                                                                                                                                                 WPI; 2003-430463/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide derived from HIV gp120 V3 loop.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUL-2003
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                                                                                                                           Claim 69; Page 136; 159pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-OCT-2001; 2001US-0341045P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UTAH ) UNIV UTAH RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 RAFVTI 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RAFVTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 28;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 9;
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The specification describes a composition for reducing an interaction between Bob and gp120. The composition comprises a Bob inhibitor that binds a region of Bob, or a substance that interacts with N-terminal

of the first loop

or the first extracellular loop domains

of Bob

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RESULT 215
ADK68776
ID ADK687
XX ADK687
XX ADK687
XX Epitop
XX epitop
XX epitop
XX cytoto
XX Virus;
OS Human
XX US2003
XX I1-DEC
XX, O7-NOV
XX (SIMA,
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PA (DI
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This invention relates to a novel method of identifying a polypeptide suitable for epitope liberation, including the steps of identifying an epitope of interest; providing substrate polypeptide sequence including the epitope, wherein the substrate permits processing by a proteasome; contacting the substrate with a composition including the proteasome; contacting the substrate with a composition including the proteasome; contacting for liberation of epitope. The invention may be useful for the development of compounds with a cytostatic, antibacterial, in controvacide or fungicide activity acting as T-cell activators. In addition, the invention may allow development of a vaccine. The invention is useful for identifying a polypeptide suitable for epitope liberation, where the epitope is a housekeeping epitope. The compositions comprising the identified housekeeping epitopes are useful in vitro in vaccine controvators. In the invention is also useful in vitro in vaccine controvators. The invention controvators is a composition of cytotoxic Tlymphocyte development or in the generation or expansion of cytotoxic Tlymphocyte (CTL) to be used in adoptive immunotherapy. The invention is also useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying polypeptide suitable for epitope e.g., housekeeping epitope, liberation by contacting substrate polypeptide comprising epitope of interest, with proteasome, and assaying for liberation of epitope.
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(DIAM/)
(QIUZ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     epitope liberation; substrate; proteasome; cytostatic; antibacterial; protozoacide; fungicide; T-cell activator; vaccine; housekeeping epitope; cytocoxic T lymphocyte; CTL; adoptive immunotherapy; neoplastic cell; virus; bacterium; protozoan; fungus; housekeeping proteasome system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        enteropathy, HIV infertility. The
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) DIAMOND D C.
) QIU Z.
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3. 1.8e+06;
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The invention describes an isolated nucleic acid (I) comprising a reading frame comprising a first sequence, where the first sequence encodes one or more segments of tumour-associated antigen SSX-2, which comprises a sequence of 188 amino acids (SEQ ID NO: 40), where the first sequence does not encode the complete SSX-2 antigen, and where each segment comprises an epitope cluster, the cluster comprising or encoding at least two amino acid sequences having a known or predicted affinity for a same MHC receptor peptide binding cleft. Also described are: an isolated polypeptide comprising the amino acid sequence encoded in the reading frame; and an immunogenic composition comprising (I) or the polypeptide of (1). (I) is a nucleic acid encoding a tumour-associated antigen SSX-2 comprising a fully defined sequence of 188 amino acids (SEQ ID NO: 40).

Disclosure; SEQ ID NO 139; 260pp; English.

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RESULT 216
ADQ10574
ID ADQ105
XX ADQ105
XX ADQ105
XX ADQ105
XX INMUNO
EN Human
XX INMUNO
KW SSX-2
KW Epitoc
KW Epitoc
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XX Human
XX INMUNO
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                    Novel nucleic acid encoding tumor-associated antigen SSX-2, useful inducing an immune response and in treating cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-517003/49
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(DIAM/)
(QIUZ/)
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07-NOV-2002; 2002US-00292413
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Pred. No.
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Best Local (
                                                                                     The peptide elicits cytotoxic T lymphocyte (CTL) response at concns. of 10(-12) to 10(-6) M. It corresp. to residues 318-327 of HIV-1 strain IIIb gp. 160 envelope glycoprotein. It can be used for the treatment and/or prophylaxis of HIV infection. (Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent clashes with ongoing US granted patent numbers. For further information please visit the Derwent web site at www.derwent.com/dwpi/updates/ntis us.html.) (Updated on 25-MAR-2003 to correct PF field.) (Updated on 27-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-AUG-2003
25-MAR-2003
17-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytotoxic T lymphocyte; immunogenic peptide; V3 glycoprotein 160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The nucleic acid, the encoded antigen, and composition are useful in inducing an immune response and in treating cancer. Expression cassettes are used in vaccine vectors. This is the amino acid sequence of a T-cell epitope MHC ligand associated with methods, therapies and compositions described in the invention.
                                                                                                                                                                                                                                                                Peptide(s) for stimulation of cytotoxic T cells specific for HIV-1 which correspond to residues 318-327 of HIV-1 gp 160 envelope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence of synthetic peptide which represents immunogenic region of the V loop of HIV isolate IIIb.
                                                                 Sequence 10
                                                                                                                                                                                                                          Disclosure; Page 8; 61pp;
                                                                                                                                                                                                                                                                                                        WPI; 1993-093577/11
                                                                                                                                                                                                                                                                                                                                  Berzofsky JA,
                                                                                                                                                                                                                                                                                                                                                                                                 26-JAN-1988;
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                                                                                                                                                                embryonic antigen, an oncogene antigen, a mutated tumour-suppressor gene antigen, or a viral antigen. They can be used for the treatment of disease such as cancer, e.g. malignant melanoma or infectious disease, e.g. viral disease such as hepatitis or AIDS. Sustained antigen delivery to the lymphatic system provides for potent CTL stimulation that takes place in the milieu of the lymphoid organ, and it sustains stimulation that is necessary to keep CTL active, cytotoxic and recirculating through the body. AAY10071 to AAY10639 represent examples of peptide antigens
                                                                                                                                                                                                                                                                                                     The present invention describes a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal. The method comprises: (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal's lymphatic system to maintain the immunologic CTL response. The method can be used for the delivery of e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Inducing a cytotoxic T lymphocyte response - antigen in the lymphatic system of a mammal sCTL response, used to treat, e.g. AIDS.
                                                                                                                      Sequence 10 AA;
                                                                                                                                                                                                                                                                                          a differentiation antigen, a tumour-specific multilineage antigen,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kuendig TM,
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10-DEC-1997;
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RAFVTI 10
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RESULT 219 AAY10547

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RESULT 220
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal. The method comprises: (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal's lymphatic system to maintain the immunologic CTL response. The method can be used for the delivery of e.g. a differentiation antigen, a tumour-specific multilineage antigen, an
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 T cell epitope/MHC ligand SEQ ID NO:94.
                                                                                                                                                                                                                                                                                                         Sequence 10 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Inducing a cytotoxic T lymphocyte response - by maintaining a level of antigen in the lymphatic system of a mammal so as to provide a sustained
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10-DEC-1997;
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                                 12-MAY-1999
                                                                 AAY10164;
                                                                                                AAY10164 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTL response, used to treat, e.g. AIDS.
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97US-00988320.
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Cytotoxic T-lymphocyte response; CTL; antigen; lymphatic immunisation; tumour; infectious disease; immunotherapy; malignant melanoma; viral disease; hepatitis; AIDS.
                                                                                                                                                          Human immunodeficiency virus 1.
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WO9902183-A2

10-JUL-1998; 98WO-US014289

10-JUL-1997; 10-DEC-1997; 97CA-02209815. 97US-00988320.

(CTLI-) CTL IMMUNOTHERAPIES CORP.

Kuendig TM, Simard JJL;

WPI; 1999-120514/10

Inducing a cytotoxic T lymphocyte response. - antigen in the lymphatic system of a mammal & CTL response, used to treat, e.g. AIDS. mammal so as to provide a sustained by maintaining a level of

Disclosure; Page 26; 199pp; English.

The present invention describes a method of inducing and/or sustaining an CC immunological cytotoxic T lymphocyte (CTL) response in a mammal. The CC method comprises: (a) delivering an antigen to the mammal at a level to cinduce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal; slymphatic system to maintain the CC immunologic CTL response. The method can be used for the delivery of e.g. cd adifferentiation antigen, a tumour-specific multilineage antigen, an embryonic antigen, an oncogene antigen, a mutated tumour-suppressor gene CC embryonic antigen, an incogene antigen, a mutated tumour-suppressor gene CC disease such as cancer, e.g. malignant melanoma or infectious disease, ce.g. viral disease such as hepatitis or AIDS. Sustained antigen delivery to the lymphatic system provides for potent CTL stimulation that takes CL place in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ, and it sustains stimulation that takes complete in the milieu of the lymphoid organ. given the present invention

Sequence 10 AA;

Query Match Best Local Matches 1 RAFVII 6 Similarity 6; Conserv Conservative 100.0%; 71.8%; Score 28; DB 100.0%; Pred. No. 25; ive 0; Mismatches DB 25; <u>ب</u> °, Length 10 Indels <u>,</u> Gaps

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RAFVTI 10

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AAY03691 standard; peptide; 10 AA

17-OCT-2003 07-JUN-1999

(revised)
(first entry)

Amino acid fragment of CTL epitope of HIV/SIV (H) string

RESULT 221
AAY03691
ID AAY036
XX AAY036
XX 17-OCT
DT 17-OCT
DT 07-JUN
XX AMINO
XX W CD8+ 7
KW CD8+ 7
KW cytotc
XX malari
XX melanc CD8+ T-cell; immune response; antigen; priming composition; CTL; epitope; cytotoxic T lymphocyte; boosting; poxvirus vector; PVV; pathogen; tumour; malaria; parasite; P. falciparum; viral; bacterial; parasitic; cancer; melanoma; HIV; breast; colon; vaccination.

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AAY03655
ID AAYC
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OT 07-X
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DE HIV
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CD8
KW CD8
KW CYC
KW CYC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CDB+ T-cell immune response against at least one target antigen. The kits of the invention comprises (i) as priming composition, a source of one or cDB+ T-cell [cytotoxic T lymphocytes-(CTL)] epitopes of the target antigen, plus a carrier and (ii) as boosting composition a source of one or complete, with at least one CTL epitope the same as used in (i), with the source being a non-replicating or replication-impaired recombinant compositions of cTL poxvirus vector (PVV) plus a carrier. If the source of CTL pitopes in (i) is a viral vector, then the vector in (ii) is from a different virus. The kits are used to generate an immune response (prophylactic or therapeutic) against pathogens or tumours, specifically against malaria parasites such as P. falciparum, or HIV, and also many other bacterial, viral or parasitic pathogens. The kits are also used for protective response against melanoma and cancer of breast or colon, and generally wherever a strong CDB+ response is protective. The boosting composition may be used alone to boost a naturally primed response against malaria. The specified PVV provide an excellent booster effect, better than that from wild-type poxvirus, resulting in complete rather than partial protection against sporozoite challenge. Also PVV are safer to use than wild-type virus. Sequences AAY0381-704 represent CTL peptide epitopes of the HIV/SIV (H) enitron entring (Hnadred on 17,007-2013 to generals).
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Matches 6
                            CD8+ T-cell; immune response; antigen; priming composition; CTL; ecytotoxic T lymphocyte; boosting; poxvirus vector; PVV; pathogen; malaria; parasite; P. falciparum; viral; bacterial; parasitic; carmelanoma; HIV; breast; colon; vaccination; PIA tumour antigen.
   Synthetic
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Hanke T, Smith
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the HIV/SIV (H)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Generating CD8-positive T cell response to target antigen recombinant poxvirus - for treating or preventing malaria infection, also epitope strings from Plasmodium and HIV.
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cive 0; Mismatches
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                                                     cancer;
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The invention relates to methods and reagents for generating a protective CC CD8+ T-cell immune response against at least one target antigen. The kits CC of the invention comprises (i) as priming composition, a source of one or CC more CD8+ T-cell (cytotoxic T lymphocytes-(CTL)) epitopes of the target CC antigen, plus a carrier and (ii) as boosting composition a source of CTL cepitopes, with at least one CTL epitope the same as used in (i), with CC this source being a non-replicating or replication-impaired recombinant CC poxvirus vector (PVV) plus a carrier. If the source of CTL epitopes in CC (i) is a viral vector, then the vector in (ii) is from a different virus. The kits are used to generate an immune response (prophylactic or CC therapeutic) against pathogens or tumours, specifically against malaria CC parasites such as P. falciparum, or HTV, and also many other bacterial, CC viral or parasitic pathogens. The kits are also used for protective compared and cancer of breast or colon, and generally compose against melaria. The specified PVV provide an excellent booster effect, better than that CC from wild-type poxvirus, resulting in complete rather than partial CC wild-type virus. SEquences AAV03653-60 represent CTL peptide epitopes coused during the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Generating CD8-positive T cell response to target antigen recombinant poxvirus - for treating or preventing malaria
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 22; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-070325/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mcmichael AJ, F
Hanke T, Smith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            infection, also epitope strings from Plasmodium and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9856919-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS INNOVATION LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-DEC-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hill
h GL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97GB-00011957
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AVS, Gilbert SC, Blanchard T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Schneider J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Plebanski M;
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prisu
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Sequence 10
Α,
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                          Query Match
Best Local S
Matches 6
ഗ
             Ь
        RAFVTI 6
                          6;
                                 Similarity
RAFVTI 10
                         ilarity 100.0%;
Conservative (
                                       71.8%;
                          0
                        Score 28; DB; Pred. No. 25; 0; Mismatches
                                 25;
                                      2
                         0,
                                      Length 10
                         Indels
                         0
                        Gaps
                         0
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AAY05357 standard; peptide; 10 A

AAY05357;

17-OCT-2003 29-JUN-1999 (revised) (first en entry)

CLUVAC peptide, SEQ ID NO. 16.

RESULT 223
AAY05357
ID AAY0535
XX AAY053
XX AAY053
XX AAY053
XX AAY053
XX IT-OCT
DT 17-OCT
DT 29-UUN
XX IT-UP
DE HIV-1
XX HIV-1
XX HIV-1;
XW PYOTEC
XW PYOTEC
XW Listes
XW immune
XX immune HIV-1; CLUVAC; cluster peptide vaccine construct; cytotoxic T lymphocyte; protective mucosal CTL response; hepatitis A virus; papilloma virus; feline immunodeficiency virus; feline leukaemia virus; M. tuberculosis; Listeria monocytogenes; M. leprae; Giardia lamblia; response induction.

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RESULT 224
AAY59593
밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents a HIV-1 cluster peptide vaccine conjugate (CLUVAC) sequence. The invention relates to a method for inducing a protective mucosal cytotoxic Tlymphocyte (CTL) response in a mammalian subject, which comprises contacting a mucosal tissue of the subject with a composition comprising a purified soluble antigen. The method can induce a protective mucosal CTL response in a subject. The method can be used for protection against e.g. hepatitis A virus, papilloma virus, feline immunodeficiency virus, feline leukaemia virus, Listeria monocytogenes, M. tuberculosis, M. leprae, or Giardia lamblia. The method induces long-lasting protective mucosal immune responses. (Updated on 17-ocT-2003 to standardise OS field)
                                                                                                                                                                                   HIV-1; env gene; cellular immunity; virus; therapy; envelope glycoprotein; infection; immunisation; imm
                                                                                                                                                                                                                                                 12-SEP-2003
05-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10 AA;
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17-FEB-1998;
                                                                                     27-MAY-1999;
                                                                                                             19-JAN-2000.
                                                                                                                                                            Human immunodeficiency virus 1
                                                                                                                                                                                                                         HIV-1 env peptide I-10
                                                                                                                                                                                                                                                                                       AAY59593;
                                                                                                                                                                                                                                                                                                               AAY59593 standard; peptide; 10 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Method for inducing a protective mucosal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-243663/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Berzofsky JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus 1.
          (NIHE-) JAPAN HEALTH SCI FOUND (AJIN ) AJINOMOTO CO INC. (UYJE-) UNIV JEFFERSON THOMAS.
                                                            29-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (USSH ) US DEPT HEALTH & HUMAN SERVICE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-MAR-1999.
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                                                                                                                                                                                                                                                                                                                                                                              RAFVTI 10
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                                                                                                                                                                                                                                                (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Belyakov IM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-0058523P.
98US-0074894P.
                                                            98US-00087513
                                                                                     99EP-00401265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US019028
                                                                                                                                                                                                                                                                                                                                                                                                                           71.5.
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                         71.8%; Score 28; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                          25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytotoxic T lymphocyte immune
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 10;
                                                                                                                                                                                    immune response
                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Strober
                                                                                                                                                                                                                                                                                                                                                                                                                               0
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CC virus. The therapeutic composition can be used to prevent or treat CC infection. Prior art methods of immunishing patients against viruses which CC frequently mutate have resulted in chronic immune activation and high T CC cell turnover because of secondary responses induced by the V3 loop CC mutated epitopes. The full length envelope glycoprotein expressed on the CC cell surface or released from HIV infected cells can also trigger CC detrimental effects which are essential in AIDS pathogenesis. The CC composition provides antigen presential in AIDS pathogenesis. The CC composition provides antigen presenting cells (APCs) which contain the CC modified envelope glycoprotein and are resistant to antibody-dependent CC cell mediated cytotoxicity (ADCC), do not form syncytia, do not undergo apoptosis and induce cellular immunity to the virus without inducing CC apoptosis of CD4+ T cells. The composition therefore redirects immune CC responses towards the conserved epitope of the envelope glycoprotein, CC inducing cellular immunity to multiple strains of the virus. (Updated on CC 12-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunodominant epitope; and (b) induces cellular immunity to a conserved epitope of the envelope glycoprotein. The nucleic acid may be introduced into a vector DNA or a liposome and mixed with an adjuvant to prepare a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention relates to a therapeutic composition for inducing cellular immunity against a virus, which comprises a nucleic acid encoding an envelope glycoprotein of the virus which: (a) contains a modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New composition for inducing viral immunity, useful for production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kaneko Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents a fragment of the HIV-1 env protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 11; 28pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      effective against and induce cellular immunity against the HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AIH
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밁 S Matches Query Match Best Local ( σ 1 RAFVTI 6 Similarity 6; Conservative RAFVTI 10 71.8%; Score 28; 100.0%; Pred. No. 0 Mismatches DB 25; ω --<u>.</u> Length 10; Indels 0 Gaps 0

Sequence 10 AA;

AAY67361 standard; peptide; 10 \$

AAY67361;

25-APR-2000 (first entry)

Human immunodeficiency virus-10 (HIV-10) peptide.

ARSULT 225
AAX67361
ID AAX673
XX AAX673
XX AAX673
XX Therap
DE Human
XX Therap
KW Tellul
KW melano
OS Human
XX WO9963
XX Human
XX Therap
PN WO9963
XX Therap
PN WO9963
XX Human
XX Therap
Therapeutic antigen; cytotoxic T lymphocyte; CTL; CTL immune response; cellular immune response induction method; vaccine; human; tumour; melanoma glycoprotein immune response induction glycoprotein 75.

Human immunodeficiency virus

WO9963945-A2

16-DEC-1999

11-JUN-1999; 99WO-US013146

12-JUN-1998; 30-OCT-1998; 98US-0089055P. 98US-0106339P.

(SLOK ) SLOAN KETTERING INST CANCER RES

Nikolic-Zugic Ģ Dyall R, Houghton AN

WPI; 2000-126432/11

Induction of a cellular immune response to a weakly immunogenic protein, used to target and kill tumor cells.

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THE SECOND SECON
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AAY94398
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; phage display; anti-inflammatory; antibody therapy; inflammatory bowel disease; rheumatoid arthritis; septic shock; multiple sclerosis; chronic inflammation; allograft rejection; tumour necrosis factor alpha; TNF; CDR3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 subject comprises administering antigen to induce a cellular immune response to the target peptide. The antigen comprises an immune portion having a major histocompatibility complex (MHC) binding domain which binds to the MHC and an immune recognition domain which is cells. The antigen is derived from the target peptide such that the MHC-binding portion binds to MHC with a greater affinity than the target peptide without material alteration of the immune response to a non-immunogenic or weakly immunogenic target peptide expressed on tumour cells of a mammalian subject. The antigens and immunogens of the invention, as well as polynucleotides encoding them,
                                                                   Small functional units of antibody heavy chain variable regions useful
                                                                                                                                            WPI; 2000-387610/33
                                                                                                                                                                                                                                                                                                                    18-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAY-2000.
                                                                                                                                                                                                         Plaksin
                                                                                                                                                                                                                                                                                                                                                                              02-NOV-1999;
                                                                                                                                                                                                                                                                (PEPT-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200029004-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV peptide used to generate a mouse hybridoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY94398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY94398 standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents a human immunodeficiency virus (HIV-10) peptide used in the method of the invention. The invention relates to a method for inducing a cytotoxic T lymphocyte (CTL) immune response to non/weakly-immunogenic proteins which are expressed on tumour cells. The method for inducing a cellular immune response to a non-immunogenic or weakly immunogenic target peptide expressed on tumour cells of a mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 15;
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                                                                                                                                                                                                                                                                PEPTOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RAFVTI 6
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K
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                                                                                                                                                                                                                                                                                                                       98IL-00127127
                                                                                                                                                                                                                                                                                                                                                                              99WO-IL000581.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     compositions against tumour cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       71.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                °,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hybridoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              panning;
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¥88888888888888888888888
                                                                      typically makes most antigen contacts in antibody combining sites. The PCR product was reamplified to avoid non-symmetric pairing of strands due to primer exhaustion. The final product was ligated into the phagemid vector pCANYAB 5 E and electroporated into E coli strain TGI. Phage clones capable of binding a specific antigen, e.g. Tumour necrosis factor alpha (TNFalpha), can be selected by library panning. Single-domain VH proteins can be used to treat or diagnose associated disorders For example, disorders in which TNF plays a role include inflammatory bowel disease, rheumatoid arthritis, septic shock, multiple sclerosis, chronic
Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                  The present sequence is an HIV peptide. A gene encoding a single-domain VH protein belonging to mouse VH group I(A) was cloned from a mouse hybridoma generated against the present sequence in complex with H-2Dd. The gene was amplified by PCR. The 3' primer contained a sequence which randomised 9 amino acids in the third hypervariable loop (CDR3) of the VH and therefore generated the single-domain VH library repertoire. CDR3
                                                         inflammation and
                                                   allograft rejection
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Matches
                                                                                                                                        Best Local Similarity
                                                                                                                                                 Query Match
 12-SEP-2003
10-JAN-2001
                             AAY94588;
                                            AAY94588 standard; peptide; 10 AA.
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                                                                                                             1 RAFVII 6
                                                                                                                              6,
                                                                                            RAFVTI 10
                                                                                                                              Conservative
(revised)
(first en
entry)
                                                                                                                                        100.0%;
                                                                                                                                                 71.8%;
                                                                                                                              0
                                                                                                                                                 Score
                                                                                                                                        Score 28;
Pred. No.
                                                                                                                              Mismatches
                                                                                                                                       DB 3;
25;
                                                                                                                              0
                                                                                                                                               Length 10
                                                                                                                              Indels
                                                                                                                            0;
                                                                                                                           Gaps
                                                                                                                             0
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밁 5

Hepatitis B virus nucleocapsid antigen; HBcAg; T cell epitope; cytolytic T lymphocyte; immunogenic; ICE; CTL; HIV; Mouse H2-d-class I restricted minimal cytolytic T lymphocyte epitope.

Human immunodeficiency virus 1.

1mmunodominant

core epitope; immunisation; mouse.

WO200026385-A1.

11-MAY-2000

05-NOV-1999; 99WO-US026291

98US-0107169P.

(POWD-) POWDERJECT VACCINES INC

Fuller DL, Fuller

2000-451623/39

Use of expression vector for nucleic acid immunization that comprises promoter and recombinant nucleic acid sequences encoding Hepatitis B ( antigen and cell epitope from antigen. core

Example 7; Page 41; 55pp; English.

RESULT 227
AAY94588
ID AAY9458
XX AAY945
XX AAY945
XX AAY945
XX HOUSE
XX HO The present invention relates to an immunogeneic recombinant nucleic a molecule. The molecule consists of a modified hepatitis B virus nucleocapsid antigen (HBcAg) with a T cell epitope sequence inserted within the HBcAg. The creation of a unique restriction site in HBcAg facilitated the insertion of the T cell epitope into the DNA encoding immunodominant core epitope of the HBcAg. An example of a suitable insertion epitope is the present sequence, the mouse H2-d-restricted acid the

Example 1; Page 18; 48pp; English

for diagnosis

and

treatment of

disease

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RESULT 228
AAB15874
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                The present invention concerns the identification of a number of chemokines which can be used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences AB15785-B15794, AB15803-B15813 and AB15831-B15848. These chemokine derivatives can be used to treat diseases such as autoimmune diseases, atherosclerosis, osteoporosis, HIV infection and AIDS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotoxaemia, myocardial infarction, acute ischaemia and theumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 minimal cytolytic T lymphocyte epitope of HIV IAI gp 120. Alternatively other T cell epitopes may be inserted (AAY94583, AAY94584, AAY94585, AAY945867). The recombinant nucleic acid molecule may then be used as a reagent in various nucleic acid immunisation strategies. The advantage of this method of immunisation is that the nucleic acid reagents that encode hybrid HBcAg generate an extremely high frequency cellular immune response against the CTL epitope. (Updated on 12-SEP-2003)
                                                                                                                                                                                                                                                                                                                                                                                                  New peptide 3, amide and heterocyclic used for inhibiting chemokine induced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               monocyte chemoattractant protein-1; inflammation; atherosclerosis; HIV; AIDS; stroke; psoriasis; autoimmune disease; hypertension; endotoxaemia; basophil-mediated disease; myocardial infarction; acute ischaemia;
                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 18; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                         stroke,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-499101/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Grainger DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NEOR-) NEORX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Macrophage recruitment; chemokine derivative; MCP-1; osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to standardise OS
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therapy
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                                                                                                                                                                                                                                                                                                                                                                           vascular diseases, autoimmune
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for the same diseases,
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99US-00452406.
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as well as in the production of
                                                                                                                                                                                                                                                                                                                                                                           compounds and saccharide conjugates activity and for treating e.g. diseases and tumor growth.
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                                                                                                                                                                                                                                                                                                                                                                         and for treating and tumor growth.
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                                               The present invention describes a modified therapeutic peptide (I) CC comprising therapeutically active amino acid region (III) and a cC reactive group (II) (e.g. succinimidyl and maleimido groups) attached to CC less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a cc peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth cf factors and neurotransmitters, to protect them from peptidase activity in cc vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptides to albumin prevents or cc modifying and attaching therapeutic peptides to albumin prevents or cc intracellular uptake and interference with physiological processes. AAB90829 to AAB9441 represent peptides which can be used in the cvents of exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local 9
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10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity.
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  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         blood component;
hydroxyl; thiol;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Virus related
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUN-2001
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99US-0153406P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Milner PG,
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Pred. No.
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides an antigenic composition comprising an antigen with a 3-O-deacylated monophosphoryl lipid A or monophosphoryl lipid A adjuvant. The presence of the adjuvant causes an increased immune response. The antigen may be from a pathogenic bacterium, fungus, virus or parasite, a cancer cell, an allergen or from amyloid peptide protein. The composition can be used in the prevention and treatment of infection, cancer and diseases caused by amyloid deposition. It is particularly useful against HIV, Neisseria gonorrhoeae and respiratory syncytial virus
Human immunodeficiency virus; HIV; immunogen; anti-HIV; vaccine; gene therapy; fusion protein; modified vaccinia virus Ankara vec
                                                                             Human immunodeficiency virus env protein derived restricted CTL epitope
                                                                                                                                              10-SEP-2001
                                                                                                                                                                                                      AAE04801;
                                                                                                                                                                                                                                                      AAE04801 standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antigenic composition having an antigen (e.g. viral protein) and an adjuvant, useful for enhancing humoral and cellular immune response host or as a prophylaxis against virus, bacterium, parasite, cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 41; 129pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB49397 standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AMCY ) AMERICAN CYANAMID CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200069456-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71.8%; 5--
100.0%; Pr
                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer; amyloid deposition
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 28; DB;; Pred. No. 25;
0; Mismatches
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25;
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  vector;
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MVA;
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Best Local Similarity
""" ches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to human immunodeficiency virus immunogens and CC their corresponding DNA molecules. An immunogen comprises a portion of CG gag protein of HIV from an HIV clade, parts of p17 and p24, modified to CG prevent N-terminal myristoylation; and a synthetic polypeptide comprising CC human cytotoxic T-lymphocyte (CTL) epitopes of HIV protein. This communogen is designed to elicit an HIV-specific immune response in CG humans. The immunogen is useful in the preparation of a medicament such cas vaccine to prevent or treat HIV infection in a human subject. The CG as vaccine to prevent or treat HIV infection in a human subject. The CG invention also relates to method of stimulating anti-HIV immune response in a human subject which comprises administering one or more times an amount of nucleic acid molecule sufficient to prime an immune response to the immunogen, or else may be packaged within a delivery means, such as a CC common portion of the immunogens. The pregent sequence is human CC immunodeficiency virus env protein derived restricted CTL epitope related to the invention. This restricted CTL epitope is presented by a murine or common portion of the immunogens to the immunogens of the invention. This restricted CTL epitope is presented by a murine company of the immunogens 
                                                                                          HIV;
vif;
                                                                  vaccine;
                                                                                                                                                            Human MHC
                                                                                                                                                                                                       15-JUL-2002
                                                                                                                                                                                                                                                    ABP25102
                                                                                                                                                                                                                                                                                             ABP25102 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel immunogen for stimulating anti-HIV immune response, has a portion of gag protein of HIV from HIV clade, parts of p17, p24 and synthetic polypeptide comprising human cytotoxic T-lymphocyte epitopes of HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-DEC-1999; 99GB-00030294.
14-OCT-2000; 2000GB-00025234.
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(ITAI-) INT AIDS VACCINE INITIATIVE.
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                                                           HIV-1; human immunodeficiency virus; env; pol; gag; nef; vpr; vpu; tat; cytotoxic T lymphocyte; CTL; immune response; epitope; antigen; ine; HIV infection; immunisation; virucide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              RAFVTI 6
                                                                                                                                                                                                                                                                                                                                                                                                               RAFVTI 10
                                                                                                                                                     peptide binding assay peptide
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                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                               peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  71.8%; Score 28; DB 100.0%; Pred. No. 25; tive 0; Mismatches
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Homo sapiens

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RESULT 233
AAJ03832
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antigens is directed largely toward variable regions of the antigen, allowing for immune escape due to mutations. The groups for inclusion in an group-based vaccine may be selected from conserved regions of viral or tumour-associated antigens, which therefore reduces the likelihood of escape mutants. Furthermore, immunosuppressive groups that may be present in whole antigens can be avoided with the use of group-based vaccines. An additional advantage of an group-based vaccine approach is the ability to combine selected groups (CTL and HTL), and further, to modify the composition of the groups, achieving, for example, enhanced immunogenicity. Accordingly, the immune response can be modulated, as appropriate, for the target disease. Similar engineering of the response is not possible with traditional approaches. ABP11501 to ABP25412 represent peptide sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccine
peptide
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Baker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 group-based vaccines has several advantages over traditional vaccines, particularly when compared to the use of whole antigens in vaccine compositions. There is evidence that the immune response to whole
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence selected from 51 defined amino acid sequences (ABL25347 to ABP25397). (I) has virucide activity and can be used in vaccines. (I) be used for immunising subjects against HIV-1 infections. The use of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a composition (I) comprising a prepared human immunodeficiency virus-1 (HIV-1) group comprising an amino acid sequence selected from 51 defined amino acid sequences (ABL25347 to ABP25397). (I) has virucide activity and can be used in vaccines. (I) maximum present the second sequence of the second sequence (I) maximum present the second sequence (I) maximum present the sequence of the second sequence (I) maximum present the seq
                                                                                                                                                                                       Hepatitis C
antiviral.
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                                                                                  WO200121189-A1
                                                                                                                                                                                                                                                                       Hepatitis C virus epitope #3823.
                                                                                                                                                                                                                                                                                                                            02-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                  AAJ03832
                                                                                                                                                                                                                                                                                                                                                                                                                                  AAJ03832 standard; peptide; 10 AA
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                                                                                                                                  Hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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Celis E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ξ
                                                                                                                                                                                                              virus; HCV; epitope; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000WO-US027766
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 28; pred. No.
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Grey HM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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25;
                                                                                                                                                                                                                 immunogen;
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                                                                                                                                                                                                                 HLA-binding motif;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a composition comprising a prepared hepatitis C virus (HCV) epitope such as those given in AAJ0010-AAJ04121. These are derived from HCV HIA-binding motifs. They are useful in vaccines for the prevention and treatment of HCV infection in humans. The present sequence is an epitope used in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus type 1; HIV-1; adjuvant; immunomodulator; alpha-2-macroglobulin; 3-0-deacylated monophosphoryl lipidA; MPL; GM-CSF; granulocyte macrophage colony stimulating factor; immune response; vaccine; R10I peptide.
The invention relates to a composition comprising activated alpha-2-macroglobulin (alpha_2M asterisk), 3-O-deacylated monophosphoryl lin (MPL) and granulocyte macrophage colony stimulating factor (GM-CSF).
                                                   Example 2;
                                                                           A (MPL) and granulocyte macrophage colony eliciting an immune response.
                                                                                                                               WPI; 2002-269315/31.
                                                                                                                                                                                                                                 27-AUG-2001; 2001WO-US026589
                                                                                                                                                                                                                                                          28-FEB-2002
                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus type 1 (HIV-1) R10I peptide
                                                                                                                                                                                                                                                                                                                                                                                                                          29-AUG-2003
18-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAE20153 standard; peptide; 10
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                                                                                                     Use of 2-macroglobulin (2Masterisk),
                                                                                                                                                                              (UYDU-) UNIV DUKE.
                                                                                                                                                                                                       25-AUG-2000; 2000US-0227624P
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This invention relates to a method for generating a herpesvirus amplicon particle comprising transfecting a cell with a Herpes simplex virus (HSV) amplicon vector, an amplicon plasmid or a nucleic acid sequence that encodes an accessory protein. The method of the invention may have anti-HIV, cytostatic; immunomodulator; antibacterial, and antiparasitic activity. The method of the invention is useful in generating herpesvirus pupilicon particles for treating patients with cancer (e.g. chronic lymphocytic leukaemia, lymphoma, glioblastoma or lung cancer) or an infectious disease such as HIV or those caused by human papilloma virus, influenza virus, bacteria or parasite. The HSV amplicon particles or the vectors can also be useful as vaccines. Gene therapy vectors based on the herpes simplex virus exhibit a broad cellular tropism, they have the capacity to package large amounts of genetic material (which makes them useful in expressing multiple genes or gene sequences), they have a high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Federoff HJ,
Dewhurst S,
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Best Local :
replicating vaccine delivery vehicle (which comprises a non-infectious recombinant viral tubule) carrying one or more immunogens. The invention is useful for inducing an immune response, preferably anti-viral, in a subject. The administration of the vaccine is preferably followed by administering one or more virus like particles carrying an immunogen. It is also useful for administering to a patient for generating antibodies specific for one or more immunogens, such as Bluetongue virus (BTV), foot and mouth disease virus (FMDV), influenza virus and human immunodeficiency virus (HTV). The invention provides an effective means of delivering multiple peptide components representing viral/tumour antigenic groups to elicit protective immunity, which has not previously been possible. The present amino acid sequence represents one of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A vaccine, for inducing an antiviral immune response, comprises a non-replicating vaccine delivery vehicle (which comprises a non-infectious recombinant viral tubule) carrying one or more immunogens.
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RESULT 237
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Celis
                                                                                                                                                                                                  Vaccine composition for treating of infection, and/or for stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    collection (AAU96022-AAU96045) of HTV epitopes that were used in methods of the invention as immunogens. These epitopes were used construct chimeric NS1-TUBs (Virus-derived tubules) which show immunogenicity. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                 Sequence 10 AA;
                                                                                                                The present invention relates to a composition comprising at least one hepatitis B virus epitope. This can be used in the production of a vaccine for use in preventing or treating hepatitis B virus infection. The present sequence is a peptide described in the exemplification of the
                                                                                                                                                                    Disclosure; Page 196; 228pp; English.
                                                                                                                                                                                       HBV peptide epitope.
                                                                                                                                                                                                                                  WPI; 2002-643192/69
                                                                                                                                                                                                                                                                                                                     08-SEP-2000; 2000WO-US024802
                                                                                                                                                                                                                                                                                                                                          08-SEP-2000; 2000WO-US024802.
                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis B virus epitope #3652.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABJ09700 standard;
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                                                                                                                                                                                                                                                                                      (EPIM-)
                                                                                                                                                                                                                                                                                                                                                                                                                             virucide;
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Kubo RT,
                                                                                                                                                                                                                                                                                                                                                                                                                            B virus; HBV;
hepatotropic;
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                                     71.8%; 5t.
100.0%; Pr
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Frey HM, Chesnut RW;
                                                                                                                                                                                                                                                                                                                                                                                                                             epitope; vaccine; HBV infection; hepatitis;
antiinflammatory.
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                                                                                                                                                                                                  preventing hepatitis B virus (HBV) n immune response to HBV, comprises
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RESULT 238 ABU57440 ID ABU574

ABU57440 standard; peptide; 10 AA.

Query Match

Score

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Length 10;

Sequence 10

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CC antiarterioscierotic, antilicer, antitrheumatic, antiarthritic, according to the pay. The MHC molecule construct is useful as a therapputic composition in in vivo or ex vivo therapy, for treating, preventing, accomposition in in vivo or ex vivo therapy, for treating, preventing, composition in in vivo or ex vivo therapy, for treating, preventing, for composition or alleviating a disease involving MHC recognising cells or establishing a prognosis of a disease or defectiveness of a medicament against a disease, involving MHC recognising cells, e.g., chronic inflammatory bowel disease such as compositing cells, e.g., chronic inflammatory bowel disease such as compositing cells, e.g., chronic inflammatory bowel disease such as community of a disease or ulcerative colitis, sclerosis, type I disease such as community of a disease, involving MHC melanoma, renal carcinoma, breast cancer, lung cancer, cancer of the cancer, leukaemia, cutaneous lymphoma, hepatic carcinoma, colorectal cancer, bladder cancer, rejection-related disease, Graft-versus-host-crelated disease, or a viral disease associated with hepatitis, Acquired Immunodeficiency Syndrome (AIDS), measles, pox, chicken pox, rubella or charges. The MHC molecule construct is also useful for flow cytometry, chistology or cytology. The present sequence represents a peptide used to create the MHC molecule construct of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New Major Histocompatibility Complex treating, preventing, stabilizing or recognizing cells e.g., cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MHC; major histocompatability complex; human; cytostatic; anti-HIV; antiinflammatory; dermatological; antiasthmatic; antidiabetic; virucide; antiarteriosclerotic; antiulcer; antirhoumatic; antiarthritic; ALIS; antipsoriatic; immunosuppressive; inflammatory bowel disease; measles; Crohn's disease; ulcerative colitis; sclerosis; type I diabetes; pox; rheumatoid arthritis; psoriasis; atopic dermatitis; asthma; chicken pox; rheumatoid arthritis; psoriasis; atopic dermatitis; asthma; chicken pox;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     molecule construct comprising a carrier molecule to which one or more molecules are attached either directly or via one or more entities. The construct of the invention may have cycostatic, antiinflammatory, dermatological, antiasthmatic, antidiabetic, anti-HIV, virucide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to a new Major Histocompatibility Complex (MHC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Fig 37; 304pp; English.
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14-MAR-2001; 2001DK-00000441.
14-MAR-2001; 2001US-0275447P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       malignant melanoma; carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-APR-2003
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DYNAL BIOTECH ASA.
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2001US-0275470P.
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Query Match
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Matches 6; Conservat
                                                                                                                                                                                                             moderate an allergic response; or a selected antigen from a molecule or its portion representing those produced by a host in an undesired manner, amount or location so as to reduce an undesired effect, in a vertebrate host. The invention is useful for increasing the ability of an antigenic composition to prevent or treat disease characterised by amyloid deposition in a vertebrate host. The invention is useful as a vaccine. The present sequence is HIV-1 peptide, IIB. This peptide corresponds to the CTL epitope within the V3 loop of HIV-1-IIB. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antigen from a pathogenic virus, bacterium, fungus or parasite, or from a cancer or tumour cell, or from an allergen, or from a self molecule; and an combination of adjuvant comprising an aminoalkyl glucosamine phosphate compound (AGP), or its derivative or analogue, and a cytokine or lymphokine, or an agonist to it. The invention is useful for increasing the ability of an antigenic composition (enhancing immune response) containing a selected antigen from a pathogenic virus, bacterium, fungus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antigenic composition for use in enhancing immune response of antigen, has selected antigen, and combination of adjuvant comprising an aminoalkyl glucosamine phosphate compound, and cytokine or lymphokine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antigenic composition; cancer; aminoalkyl glucosamine phosphate compound; AGP; immune response; cytotoxic T lymphocyte; allergic response; tumour; amyloid deposition; vaccine; antifungal; antibacterial; antiparasitic; cytostatic; immunostimulant; virucide; HIV-1 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or parasite to elicit an immune response especially cytotoxic T lymphocytes; a selected antigen a cancer or tumour cell to elicit therapeutic or prophylactic anti-cancer effect; a selected allergen to
                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an antigenic composition comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 28; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-NOV-2000;
18-OCT-2001;
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14-NOV-2002
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                                                acquired immunodeficiency syndrome (AIDS). The nucleic acids may be used to treat a person at risk of becoming CD4+ T cell-deficient, particularly where someone at risk of becoming CD4+ T cell-deficient, particularly infection, particularly by an intracellular pathogen, especially one caused by cytomegalovirus, Mycobacterium tuberculosis, M. avium, Epstein-Barr virus, a fungus yeast, varicella zoster virus or human immunodeficiency virus (HIV). The present sequence is a HIV-1 class I-restricted gp120 peptide, used in the exemplification of the invention. (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                      cytotoxic T lymphocyte (CTL) activity in a CD4+ T cell-deficient individual, comprising administering an immunostimulatory sequence oligonucleotide (ISS-OND). The immunostimulatory mucleic acids of the invention are used in CD4+ T cell-deficient individuals to decrease tumour load, to treat a primary or acquired immunodeficiency, particularly where the acquired immunodeficiency is temporary and due cancer radiotherapy or chemotherapy or immunosuppression following bor marrow or organ transplantation, or acquired immuno disease treatment, or immunosuppression to the control of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-MAR-2000; 2000US-0192537P.
11-MAY-2000; 2000US-0203567P.
05-JUL-2000; 2000US-0215895P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunostimulatory sequence oligonucleotide; ISS-ODN; chemotherapy; immunosuppression; transplantation; autoimmune disease; infection; acquired immune deficiency syndrome; AIDS; intracellular pathogen; cytomegalovirus; mycobacterial infection; Epstein-Barr virus; varicella zoster virus; human immunodeficiency virus; HIV.
   Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Increasing antigen-specific cytotoxic T lymphocyte activity in a CD4+cell deficient individual, useful to treat immunodeficiency and block infection, comprises administering immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Raz E,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytotoxic T lymphocyte;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention relates to a method for increasing antigen-specific
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                                                   Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                   The present invention relates to a method for eliciting an immune response in a subject. The method comprises administering a peptide or protein antigen comprising T-cell epitope(s) (e.g. tumour antigen, viral or non-viral antigen) coordinately with a non-viral vector comprising a polynucleotide encoding a T-cell co-stimulatory molecule. Viral peptide antigens may include human immunodeficiency virus (HTV) antigen, hepatitis B virus (HBV), herpes simplex virus (HSV) or human papilloma virus (HBV). The method is useful to elicit an immune response in a subject, and to supplement and enhance peptide and protein based vaccines and treatment methods. ABG68840-ABG68700 represent HTV-1 peptide antigens useful in the method of the present invention. (Updated on 29-AUG-2003 to
                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 7; 39pp;
                                                                                                                                                                                                                                                                                                                                                                           Administering a non-viral vector encoding a co-stimulatory molecule alongside a peptide or protein T cell epitope, elicits increased re to the antigen and is useful to enhance peptide and protein based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KHLE/)
(BERZ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eliciting an immune response; peptide antigen; T-cell epitope; tumour antigen; viral antigen; non-viral vector; HIV-1; T-cell co-stimulatory molecule; human immunodeficiency virus;
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07-OCT-2002
                                                                                                            Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                vaccines and treatments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus 1; (IIIB isolate)
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                                                                                                                                          standardise OS field)
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BERZOFSKY J.
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RAFVTI 10
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                                                       Conservative
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tive 0; Mismatc
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RESULT 242 ABG80230 ID ABG802

ABG80230 standard; peptide; 10

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RESULT 243
ABG79846
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                                                                                                                                                                                                                                                                                                                           The invention relates to a method of inducing and/or sustaining an CC immunological cyrotoxic T lymphocyte (CTL) response in a mammal comprising administering directly to the lymphatic system of the mammal: CC comprising administering directly to the lymphatic system of the mammal: CC an antigen in the form of a polypeptide; (b) a vector comprising a nucleic acid encoding the antigen; or (c) a non-peptide antigen. The cc method is useful for inducing and/or sustaining CTL response in a mammal. CC method is particularly useful for treating a mammal having a malignant thumour (e.g. carcinoma, melanoma, leukaemia or lymphoma) or infectious confidence (e.g. hepatitle, acquired immune deficiency syndrome (AIDS), comparis, measles or tuberculosis), or in an animal having a cats, mice, cattle, sheep, pigs, goats, rabbits, or preferably humans. ABG79753-CC ABG80319 represent viral epitopes on major histocompatibility complex (MIC) class I molecules, used in the method of the invention. (Updated on CC 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Inducing or sustaining immunological cytotoxic T lymphocyte response in mammal, useful for treating a mammal with malignant tumor or infectious disease, by directly administering an antigen to the lymphatic system of
 MHC class I molecule, viral epitope #94.
                                   15-NOV-2002
                                                                                                   ABG79846 standard; peptide; 10 AA
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15-NOV-2002
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal comprising administering directly to the lymphatic system of the mammal:
(a) an antigen in the form of a polypeptide; (b) a vector comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Inducing or sustaining immunological cytotoxic T lymphocyte response in a mammal, useful for treating a mammal with malignant tumor or infectious disease, by directly administering an antigen to the lymphatic system of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Major histocompatibility complex; MHC; MHC class I molecule; virus; epitope; cytotoxic T lymphocyte response; CTL response; lymphatic system antigen; immunogenic; malignant tumour; carcinoma; melanoma; luukaemia; lymphoma; infectious disease; hepatitis; malaria; measles; tuberculosis; acquired immune deficiency syndrome; AIDS.
                                                                                  Cytolytic T lymphocyte; epitope; vaccine; prophylaxis; HIV infection; human immunodeficiency virus; acquired immune deficiency syndrome; CTL;
                                                                                                                                      HIV CTL epitope #12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (a) an antigen in the form of a polypeptide; (b) a vector comprising a nucleic acid encoding the antigen; or (c) a non-peptide antigen. The method is useful for inducing and/or sustaining CTL response in a mammal. This is particularly useful for treating a mammal having a malignant
                                Human immunodeficiency virus
                                                                 gene therapy; AIDS
                                                                                                                                                                           28-MAY-2003
                                                                                                                                                                                                                                            AAE35167 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10 AA;
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WO200294313-A2

New recombinant cell comprising an exogenously derived nucleic acid

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a recombinant nucleic acid comprising a nucleic acid sequence encoding an antigen containing two or more cytolytic T lymphocyte (CTL) epitopes or its analogues. Sequences of the invention are used in vaccines and are useful for the treatment and prophylaxis of human immunodeficiency virus (HIV) infection, particularly acquired immune deficiency syndrome (AIDS). The invention is also useful in gene therapy. The present sequence is HIV CTL epitope. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV antigenic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fuller D,
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18-MAY-2001; 2001US-0291655P.
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                                                         WPI; 2003-103265/09.
                                                                                                                  Rock KL,
                                                                                                                                                                                                                                  30-MAR-2001; 2001US-0280669P
                                                                                                                                                                                                                                                                                           01-APR-2002; 2002WO-US010385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T-cell immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOP; thimet oligopeptidase; EC3.4.25.15; cytostatic; tumour; immunostimulant; major histocompatibility complex class I; MHC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP60029 standard; peptide; 10 AA
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6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 28; DB;
Pred. No. 25;
0; Mismatches
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RESULT 246
ABR39122
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XX ABR391
XX ADP-ri
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a wild-type cell from which the recombinant cell is derived. The activity of TOP may be described as cytostatic and immunostimulatory. Thimet oligopeptidase (TOP, EC3.4.25.15) plays a key role in modulating levels of major histocompatibility complex (MHC) class I-presented peptides. The recombinant host cell of the invention is useful for modulating an antigenic response in a mammal. Methods of the invention are useful for screening a test compound for its ability to serve as an immunomodulatory agent and identifying an antigen resistant to thimet oligopeptidase degradation. A method of the invention is useful for increasing CDB T-cell immunity, which uses vaccination with a TOP inhibitor for decreasing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOP expression or activity. The vaccination method uses treated tumour cells, antigen bearing/pulsed dendritic cells or injection of a viral vector. The recombinant host cell is useful for treating tumours. The current sequence represents an HIV (human immunodeficiency virus) gpl60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          derived nucleic acid that codes for a thimet oligopeptidase (TOP) polypeptide. The TOP polypeptide is overexpressed in the cell compared to a wild-type cell from which the recombinant cell is derived. The activity
                                                                                New compositions comprising nucleic acid adjuvants, useful in immunization techniques, particularly for eliciting or enhancing immune response against an antigen in a human.
                                                                                                                                                                                                                                                                                                                                                                                                      ADP-ribosylating exotoxin; immune response; immunisation; vaccine; adjuvant; HIV; gp120; CTL epitope.
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10-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       coding for a thimet oligopeptidase polypeptide, useful for modulating an antigenic response in a mammal for treating e.g., tumor.
                                                                                                                                                WPI; 2003-221541/21
                                                                                                                                                                                                                                              27-NOV-2000; 2000US-00724315
                                                                                                                                                                                                                                                                            26-NOV-2001; 2001WO-US043151.
                                                                                                                                                                                                                                                                                                             16-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus 1.
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                                                                                                                                                                                                              (POWD-) POWDERJECT VACCINES INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTL epitope peptide SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
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100.0%; Pr
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Pred. No.
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The invention relates to a composition comprising: acid sequence that is a truncated A subunit coding

(a) a first nucleic
region obtained or

The present sequence is that of an HIV-specific antigen comprising the CD8+ T cell epitope of HIV-1 pl8 protein. A claimed method for enhancing a T cell response to HIV antigen in a susceptible mammalian (human) host comprising co-administering an HIV-specific antigen such as the present antigen and alpha-galalctosylceramide as adjuvant. The HIV-specific antigen may be presented by a recombinant virus such as a recombinant adenovirus, pox virus or Sindbis virus. This is an example of the method of the invention for augmenting the immunogenicity of an antigen in a mammal by immunising the mammal with the antigen and with a glycosylceramide adjuvant. This adjuvant enhances and/or prolongs antigen

55; Page 75; 97pp;

English.

English.

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RESULT 247
ABP72314
ID ABP723
XX ABP723
XX ABP723
XX ABP723
XX HIV-1
CX HUWAN
CF HUW
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                                                                                                                                                                                                                                                                                                                                                                                                                  Augmenting the immunogenicity of an antigen in a mammal, useful for treating cancer, viral infection and malaria, comprises immunizing mammal with the antigen conjointly with adjuvant comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-268011/26.
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08-MAY-2003
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6; Conserv
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(first entry)
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                                                                                                                      This invention relates to a novel DNA expression vector for inducing an immune response. The DNA expression vector of the invention encodes both an immunopotentiating chemokine sequence as well as an immunopotent concerous cell. The chemokines are preferably selected from the animal to cancerous cell. The chemokines are preferably selected from the animal to be treated. The vaccine of the invention may have immunostimulant, immunosuppressant and cytostatic activities and used for a form of gene therapy. The expression vector and compositions comprising the vector of the invention may therefore be useful for inducing an immune response in a mammal, and for treating cancers (tumours and metastatic form of cancer), infectious diseases, autolimiune diseases and other diseases that
cancer), infectious diseases, autoimmune diseases and other diseases the can be alleviated by either stimulating or suppressing T cell activity. The present sequence is the amino acid sequence of the human immunodeficiency virus gp120 peptide which was used during the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New DNA expression vectors comprising a DNA encoding an immunopotentiating chemokine and a DNA encoding a heteroimmunogenic polypeptide, useful for inducing an immune:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA expression vector; immune response; immunopotentiating chemokine; immunogenic polypeptide; infectious agent; cancerous cell; immunostimulant; immunosuppressant; cytostatic; gene therapy; cancer; tumour; metastatic cancer; infectious disease; autoimmune disease; stimulating T cell activity; suppressing T cell activity; macrophage inflammatory protein; HIV; gp120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -specific Th1-type responses, particularly CD8+ T cell responses. In an example from the invention, co-administration of alpha-galactosylceramide to mice immunised with the present T cell epitope enhanced almost 3 times the level of HIV-specific CD8+ T cell response compared to that induced in mice immunised without alpha-galactosylceramide treatment. (Updated on 23-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Generating an immune response against a non-hepadnaviral antigen in a mammal, useful for treating or preventing cancer or malaria, by administering a priming component comprising a recombinant hepatitis B
                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                            represents a peptide from human immunodeficiency virus type used as a carrier peptide to augment the immune response aga
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                                                                                                                                                                                                     Sequence 10 AA;
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non-hepadnaviral antigen; hepatitis B core particle; CD8+ T-cell;
epitope; poxvirus vector; cancer; malaria; epitope.
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Match
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Listing first)
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hits satisfying chosen parameters:
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PCT-US94-02631-52
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US-08-704-704-70-74
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US-08-704-704-70-74
PCT-US94-02631-73

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US-08-480-332-3

Sequence 3, Application US/08480332

; Patent NO. 6180134

; GENERAL INFORMATION:
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Patent No. 5622933
GENERAL INFORMATION:
                                                                                                                                                         Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/260,086
FILING DATE: 15-JUN-1994
CLASSIFICATION: 514
FRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9318901.7
FILING DATE: 13-SEP-1993
ATTORNEY/AGENT INFORMATION:
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248
249
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INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: WEIL, GOTSHAL & MANGES
STREET: 2882 SAND HILL ROAD
CITY: MENLO PARK
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: 37965.0007
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 926-6200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                             NAME: RAE-VENTER, BARBARA
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                                                                                                                              RAFVIIGK 8
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GLUCKMAN, JEAN-CLAUDE
VAN RIETSCHOTEN, JURPHAAS
                                                                                                                                                         100.0%; Score 39; DB 1; llarity 100.0%; Pred. No. 4.1e+05 Conservative 0; Mismatches
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BENJOUAD, ABDELAI
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                                                                                                                                                                         4.1e+05;
                                                                                                                                                                                      Length 8;
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,332
FILING DATE: 7-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/316,436
FILING DATE: 29-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/316,436
FILING DATE: 29-SEP-1994
PRIOR APPLICATION NUMBER: US/08/035,443
APTIORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 38,563
REFERENCE/DOCKET NUMBER: 5325-0115.31
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
                                                                                                                                                                                                    GENERAL INFORMATION:
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Best Local Similarity
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TELEFAX: (415) 324-0960
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY:
LOCATION:
                                                                                                                                        APPLICANT:
APPLICANT:
              APPLICANT: SASTRY, Jagannadha K.
APPLICANT: ARLINGHAUS, Ralph B.
APPLICANT: PLATSCOUCAS, Chris D.
APPLICANT: NEHETE, Pramod N.
TITLE OF INVENTION: METHODS AND COMPOSITIONS
TITLE OF INVENTION: FOR ELICITING IMMUNE OR ANTI-INFECTIVE RESPONSES
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS LENGTH: 8 amino acids
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MEDIUM TYPE: Floppy disk
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APPLICANT: Barenholz, Yechezkel
TITLE OF INVENTION: Enhanced Circulation Effector Composition and
TITLE OF INVENTION: Method
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSE: Dehlinger & Associates
STREET: 350 Cambridge Avenue, Suite 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORIGINAL SOURCE:
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HYPOTHETICAL:
CORRESPONDENCE ADDRESS:
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STRANDEDNESS: unl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide 3, Fig. 13
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ADDRESSEE: Arr STREET: P.O. E CITY: Houston

E: Arnold, White & Durkee P.O. Box 4433

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RESULT 4
US-08-704-170-44
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MOLECULE TYPE: peptide
PCT-US92-10378-5
                                                                                                                                                                                                                                                                                                                                                                     Patent No. 5707626
GENERAL INFORMATION:
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Best Local (
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TELEX: Not Applicable
INFORMATION FOR SEQ ID NO:
           COMPUTER READABLE FORM:

MEDIUM TYPE: Ploppy disk

COMPUTER: IEM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/704,170
                                                                                                                                                                           NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSE: Robbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite 500
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/945865
FILING DATE: September 16, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 512-320-7200
TELEFAX: 512-474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 07/800,93
FILING DATE: December 2, 1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
07/800,932
FILING DATE:
                                                                                                                                                           COUNTRY: U.S.A.
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STRANDEDNESS: Sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Parker, David L.
REGISTRATION NUMBER: 32
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 199212
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                                                                                                                                             90012
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5707626
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8; Conserv
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         US/08/704,170
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Pred. No. 4.1e+05;
; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/09/454,204A
CURRENT FILING DATE: 1999-12-09
PRIOR APPLICATION NUMBER: PCT/GB98/01681
PRIOR FILING DATE: 1998-06-09
PRIOR APPLICATION NUMBER: GB 97 11957.2
PRIOR FILING DATE: 1997-06-09
NUMBER OF SEQ ID NOS: 78
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 52
                                                                                                                                LENGTH: 9
; TYPE: PRT
; ORGANISM: Unknown
; FEATURE;
; OTHER INFORMATION: CTL Epitope of HIV-1 gp120
US-09-454-204A-52
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US-09-454-204A-52
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                                       Query Match
Best Local Similarity 100
Ches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; MOLECULE TYPE: peptide US-08-704-170-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
TELEPAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 52, App...
No. 6663871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: McMichael, Andrew
APPLICANT: Hill, Adrian V.S.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Schneider, Jorg
APPLICANT: Plebanski, Magdalena
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Methods and Reagents for Vaccination TITLE OF INVENTION: Which Generate A CD8 T Cell Immune FILE REFERENCE: 2907.1000-000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 9 amino a TYPE: amino acid TOPOLOGY: linear
2 RAFVTICK 9
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                             RAFYTIGK 8
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Smith, Geoffrey L.
Blanchard, Tom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09454204A
                                                                 100.0%; Score 39; DI
100.0%; Pred. No. 4.1
tive 0; Mismatches
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                                                                                     4.1e+05;
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                                                                                                     Length 9;
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RESULT 6
PCT-US94-02631-44
; Sequence 44, Application PC/TUS9402631
; GENERAL INFORMATION:

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US-08-648-298-4
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                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Henrik
APPLICANT: Eric Pa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 201 ...
STREET: Los Angeles
CITY: Los Angeles
CTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 19:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 01
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                               NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: Darby & Darby PC
STREET: 805 Third Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
           SOPTWARE: PatentIn Release #1.0, Version #1.30 (EPO) CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/648,298
                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                  STATE: Ni
STATE: Ni
10022
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                                                             MEDIUM TYPB: Diskette
COMPUTER: IBM PC compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 y Match 100.0%;
Local Similarity 100.0%;
nes 8; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE:
                                                                                                                                                                               CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH:
                                                                                                                                                                                                                                                                  LE OF INVENTION: UDP-N-acetyl-alpha-D-galactosamine:polypeptide OF INVENTION: N-acetylgalactosaminyltransferase GalNAc-T3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 RAFVTICK 8
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201 North Figueroa Street, Suite 500
                                                                                                                                                                                                                                                                                                      Henrik Clausen
Eric Paul Bennett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ehresmann,
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15-JUN-1996
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Pred. No. 4.1e+0
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                    FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
PILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 236687
INFORMATION FOR SEQ ID NO:
                                                                    TELEFAX: (213) 977-10
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: HIV gp120
                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPACTION
OFFRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                               SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                 NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Green, Reza
REGISTRATION NUMBER: 38,475
                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 201 No. 57
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
TOPOLOGY:
                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                    TELEPHONE:
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Takehana, Yoshi
Ehresmann, Glenn
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                                                                                                    (213) 977-1001
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                                                                                   977-1003
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                                                                                                                                                                                                                             US 08/029,850
                                                                                                                                                                                                                                                                                                  US/08/704,170
                                                                    52:
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                                                                                                                                         1920-331
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                                                            Best Loc
Matches
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                                                                                             Query Match
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Best Local Similarity
Matches 8; Conservat
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                                                                                                                                                                                                   TELEX: 421792
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08\326,676

FILING DATE: 07\726,605

APPLICATION NUMBER: 07\726,605

FILING DATE: 09-July-1991

APPLICATION NUMBER: 07\663,262

FILING DATE: 01-Mar-1991

APPLICATION NUMBER: 07\155,321

FILING DATE: 12-Feb-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Chang Yi Wang
TITLE OF INVENTION: SYNTHE
TITLE OF INVENTION: HUMAN
TITLE OF INVENTION: HUMAN
TITLE OF INVENTION: AND AS
NUMBER OF SEQUENCES: 38
                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 212-758-4800
                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Maria C. H. Lin
REGISTRATION NUMBER: 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                            Local Similarity
nes 8; Conserv
                                                                                                                                                                                                                                             TELEPHONE: 212-,55
TELEPHONE: 212) 751-6849
                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                      TYPE: Amino acids
                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 1151-4004 US4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION:
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                           RAFVTICK 8
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                                                                                                                                                                                           12 amino acids
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345 PARK AVE.
                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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                                                                                                                                            Unknown
                                                     100.0%; Score 39; DB 1; Length 12; 100.0%; Pred. No. 0.058; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SYNTHETIC PEPTIDES AND PROCESS
OF USING SAME FOR THE DETECTION OF ANTIBODIES TO
HUMAN IMMUNODEFICIENCY VIRUS (HIV) GP120 ENVELOPE
PROTEIN, DIAGNOSIS OF AIDS AND PRE-AIDS CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AND AS VACCINES
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Pred. No. 0.058;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>.</u>
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                                                     Gaps
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Best Local S
Matches 8
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PCT-US94-02631-52
                                                                                                                                                                                       Sequence 43, Application PC/TUS9503236
GENERAL INFORMATION:
APPLICANT: University of Southern California
TITLE OF INVENTION: Methods to Diagnose and 7
TITLE OF INVENTION: Infection
NUMBER OF SEQUENCES: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 52, Application:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (213) 977-10
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,21
REFERENCE/DOCKET NUMBER: 1
            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
TELEFAX: (213) 977-1003
FORMATION FOR SEQ ID NO: 52:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                              STREET: 4370 La Jolla Village Drive, Suite
CITY: San Diego
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Takehana,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Douvas, Angeline
                                                                                   COUNTRY:
                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity nes 8; Conserv
 OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR ITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                       1 RAFVTICK 8
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California
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201 North Figueroa
                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ehresmann, Glenn
                                                                                                                                                        Campbell and Flores
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide
PC-DOS/MS-DOS
                                                                                                                                                                                                                         Methods to Diagnose and Treat HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
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igueroa Street, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29,215
                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                         .058;
                                                                                                                                     700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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US-08-090-148-5
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                         REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELESPHONE: (212) 940-8636
TELEFAX: (212) 940-6404
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 AMINO ACIDS
TYDE: MATERIAL STATES
TYPE: MATERIAL STATES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Mastico
APPLICANT: Stockle
APPLICANT: Talbot,
                                                                                                                                                                 APPLICATION NUMBER: US/08/090,148
FILING DATE: 08/11/93
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9101550.3
FILING DATE: 01/24/91
APPLICATION NUMBER: PCT/GB92/00124
FILING DATE: 01/22/92
ATTORNEY/AGENT INFORMATION:
NAME: Nissenbaum, Israel
REGISTRATION NUMBER: 27,582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Imbra, Richard J.
REGISTRATION NUMBER: 37,643
REFERENCE/DOCKET NUMBER: FP-S
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      equence 5, Application US/08090148 atent No. 5534257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Rosenman & Colin
STREET: 575 Madison Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   y Match 100.0%; Score 39; DB 5; Local Similarity 100.0%; Pred. No. 0.058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PS2-486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10022-2585
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                 AMINO ACID
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Stockley, Peter George
Talbot, Simon John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PatentIn Release #1.0, Version #1.25
NOT RELEVANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diskette, 3.5", 1.44Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antigen-Presenting Capsid with 
Pusion MS2-Coat Protein
                                                                                                                                                                                                                                                                                                                                                                                                                          MS-DOS
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                                                                                                                                                                                                                                                PCT/GB92/00124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 5; Length 12;
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RESULT 14
US-08-111-080-6
                                                                                                                                                                                                             US-08-279-906A-17
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                                                                                                                                       Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: NO. 5619922and, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 32028
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-0448
                                                                                                                                                                                                                                                                                                                 TELEFAX: (312) 474-04-
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: Chicago
CITY: Chicago
Illinois
                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS: ADDRESSE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Borun
                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Terada, Masaki
APPLICANT: Yoneda, Yukio
TITLE OF INVENTION: NM03 Antibody Materials and Methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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Local Similarity 100.0%; Score 39; DB 1;
Local Similarity 100.0%; Pred. No. 0.063;
hes 8; Conservative 0; Mismatches o
                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 60606
                                                                                                      1 RAFVTICK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7, Application US/08279906A
5618922
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                                                                     RAFVTICK 13
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                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ohno, Tsuneya
                                                                                                                                                       100.0%; Score 39; DB 1; Length 13; 100.0%; Pred. No. 0.063;
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Sequence 6, Application 08/111080 Patent No. 5558865
GENERAL INFORMATION:
APPLICANT: Ohno, Tsuneya

APPLICANT: Ohno, Tsuneya
TITLE OF INVENTION: HIV Immunotherapeutics
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:

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RESULT 15
US-08-211-980-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; MOLECULE TYPE: US-08-111-080-6
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Best Local (
                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Ohno, Tsuneya
TITLE OF INVENTION: HIV
                                                                                                                               STREET: 6500
CITY: Chicago
CTATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
               SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 3
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APPLICATION UMBER: PCT/US92/07111
PILING DATE: 24-AUG-1992
PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: BAPLICATION DATA: US 08/039.457
FILING DATE: US 08/039.457
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
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 APPLICATION NUMBER:
                                                                                                                     COUNTRY: U. ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (312) 474-0448
                                                                                                                                                                                                   ADDRESSEE: Marshall, O'Toole, Gerstein, Murray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 14 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Borun, Michael F. REGISTRATION NUMBER: 25,447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
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STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE:
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8; Conserv
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6300 Sears Tower,
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22-APR-1993
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US/08/211,980
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                                                                                                                                                                                 Wacker Drive
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Patent No.
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SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PAPPLICATION NUM
                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERRNCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-WAX-1993
ATTORNEY/AGENT INFORMATION:
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APPLICANT:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITI
TITLE OF INVENTION: ELICITING
TITLE OF INVENTION: LYMPHOCY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Palan-
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REGISTRATION NUMBER: 25,447
REFERENCE/DOCKET NUMBER: 316
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 24-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
tes 8; Conserv
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TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OF SEQUENCES:
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5932218
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ahlers, Jeffrey D. Pendleton, C. D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Berzofsky,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (312) 474-0448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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                                                                     1173-434P
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Pred. No. 0.068;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Version #1.25
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Best Local Similarity
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                                                                                                                          TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acide
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                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
FILING DATE:
                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 14-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                 TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REPLICANT: Shirai, mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: Peptide LOCATION: 1..14 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
                                                                                                                                                                                               TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 22040-0747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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GY: linear
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                                                                                                                            14 amino acids
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Ahlers, Jeffrey D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= peptide
/note= "p18-1, see Table
                                                                                                                                                                                                                                                                                                                                US 08/060,988
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Query Match
Best Local Similarity
""" hes 8; Conserve
                                                                                       US-08-455-685-9
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US-08-455-685-9
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Best Local
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAX-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/060,988
APPLICATION NUMBER: 08/060,988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9,
                                                                                                                                                                                                                                                                                                                         FILING DATE: 14 MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06 MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                        TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPH
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                              TELEFAX: 51.
                                                                                                                                                                                                                                                                            NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022003
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OTHER INFORMATION:
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ZIP: 02110-2804
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                                                                                                                                                                                                                             617/542-8906
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                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jeffrey D.
                                100.0%; Score 39; DB 3; 100.0%; Pred. No. 0.068;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /label= peptide
/note= "pl8-2, see Table V"
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                  Mismatches
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                                               Length 14;
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                  Indels
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1 RAFVTIGK 8

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                                  Sequence 9, Application US/08060988A Patent No. 6294322
GENERAL INFORMATION:
                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                               Query Match
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TELEX: 200154
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
ENCOTH: 14 amino acids
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APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETER
TITLE OF INVENTION: MEUTRALIZI
TITLE OF INVENTION: MEUTRALIZI
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REGISTION NUMBER: P-42,3
REFERENCE/DOCKET NUMBER: 08
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08
FILING DATE: 31-MAY-1995
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
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APPLICATION NUMBER:
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02110-2804
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Berzofsky, Jay A.
Ahlers, Jeffrey D.
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Ahlers, Jeffrey D.
Pendleton, C. David
                                                                                                                                                                                                                                                                                                                                    linear
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NUMBER: 07/148,692
26-JAN-1988
                                                                                                                                                                                                                                                                                                                peptide
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                                                                                                                                                                                                                          100.0%; S
100.0%; E
ative 0;
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HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FAST-SEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 39; DB 3; Length 14; Best Local Similarity 100.0%; Pred. No. 0.068; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INCOMPATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino
                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                      APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                   APPLICANT:
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APPLICANT: Nara, Pet
APPLICANT: Shirai, M
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5970
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                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
TOPOLOGY: linear
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                                                               OF SEQUENCES:
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: Fish & Richardson P.C.
225 Franklin Street
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617/542-8906
                                                                                                                                                                              Nara, Peter
                                                                                                                                                                                               Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. David
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VENTION: MULTIDETERMINANT PEPTIDES
VENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
VENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pendleton, C. David
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application PC/TUS9207111
GENERAL INFORMATION:
APPLICANT: Ohno, Tsuneya
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CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/748,562
PILLING DATE: 22-AUG-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FABESEQ for Windows Version CURRENT APPLICATION DATA:
APPLICATION NUMBER.
                                                                                        SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: PCT/US92/07111
FILING DATE: 19920824
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FILING DATE: 14-MAY-
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: HIV Immunotherapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 08
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                            MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 100.0%; Score 39; DB 3; Length 14; Local Similarity 100.0%; Pred. No. 0.068; Indels : 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/847,311 FILING DATE: 06-MAR-1992 APPLICATION NUMBER: 07/751,998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Beattle, Ingrid A. REGISTRATION NUMBER: P-42,306
                                                                                                                                                                                                                                                                            CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/1
                                                                                                                                                                                                                                             ZIP: 60603
                                                                                                                                                                                                                                                           COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Bicknell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 RAPVTICK 14
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14-MAY-1993
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PCT-US93-07967-6
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PCT-US93-07967-6
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GENERAL INFORMATION:
APPLICATT: Ohno, Tsuneya
TITLE OF INVENTION: HIV Immunotherapeutics
NUMBER OF SEQUENCES: 38
Query Match 100.0%; Score 39; DB 5; Best Local Similarity 100.0%; Pred. No. 0.068; Matches 8; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (312) 984-9740
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
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Best Local Similarity
                                                                                                                                                                                   TELEFAX: (312) 474-04-
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                      NAME: Borun, Michael F.
REGISTRATION NUMBER: 25,447
REFERENCE/DOCKET NUMBER: 310
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: Chicago
CITY: Chicago
Illinois
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REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 31016
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 346-5750
TELEPAX: (312) 984-9740
                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 00 FILING DATE: 22-APR-1993 ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: PCT/US92/07111
FILING DATE: 24-AUG-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                     TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-0448
                                                                                                               TOPOLOGY:
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CLASSIFICATION:
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TYPE: AMINO ACID
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ADDRESSEE: Borun
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                                    DB 5;
                                  Length 14;
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   Indels
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CORRESPONDENCE ADDRESS:

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Query Match
Best Local Similarity
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PCT-US94-05142-10
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PCT-US94-05142-9
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                                                                                      Sequence 10, Application PC/TUS9405142 GENERAL INFORMATION:
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 703-205-8050
NFORMATION FOR SEQ ID NO:
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MOLECULE TYPE:
FRAGMENT TYPE:
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TITLE OF INVENTION: I
TITLE OF INVENTION: I
NUMBER OF SEQUENCES:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 0
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: Peptide LOCATION: 1..14 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330 REFERENCE/DOCKET NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 13-MA' CLASSIFICATION:
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T: P.O. Box 747
Falls Church
Virginia
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internal
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                                                                                                                                                                                                                                                                        100.0%; Score 39; DB 5; Length 14; 100.0%; Pred. No. 0.068;
                COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC LYMPHOCYTES AGAINST HIV
                                                                                                                                                                                                                                                                                                                                   /label= peptide
/note= "p18-1, see Table V"
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Patent No.
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Best Local S
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                        COUNTRY: USA
ZIP: 20005-3315
COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                       ADDRESSEE: punner
ADDRESSEE: Dunner
ADDRESSEE: Dunner
ADDRESSE: 1300 I Street, N.W.
                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                 NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                      APPLICANT: Ullmann, Agnes
TITLE OF INVENTION: Specific Immune Responses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION UNMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Peptide LOCATION: 1..14 OTHER INFORMATION: OTHER INFORMATION:
COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: P.O. Box 7. CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
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8; Conserv
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O. Box 747
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internal
PC-DOS/MS-DOS
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/note= "p18-2, see Table V"
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US-08-336-087-2
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                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PatentIn Release #1.0, Version #1:25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/218,025A
FILING DATE: 24-MAR-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA:
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                                                                            TELEPHONE: (215) 540-581
TELEFAX: (215) 540-581
INFORMATION FOR SEQ ID NO:
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INFORMATION FOR SEQ ID NO:
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SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
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PRIOR APPLICATION UNMBER: US
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/336,087
                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 540-9206 TELEFAX: (215) 540-5818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                         NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: WST33A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT: Weiner, David B.
PPLICANT: Ugen, Kenneth E.
PPLICANT: Williams, William V.
ITLE OF INVENTION: Methods and Compositions for Diagnosing
ITLE OF INVENTION: and Treating Certain HIV Infected Patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 03495-0109-01000
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STATE: Pennsylvania
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TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
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8; Conserv
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P.O. Box 457, 321 No. 5556744ristown Road
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US-08-709-047-7
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                                                                                         US-08-479-400-2
                                                                                                       RESULT 29
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                                        Sequence 2, Application US/08479400 Patent No. 5679784 GENERAL INFORMATION:
                                                                                                                                                                                                              Matches
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Best Local S
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APPLICANT: Fung, N
APPLICANT: Liming
                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids
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MEDIUM TYPE: Diskette, 3.5 i
COMPUTER: IBM PS/2
OPERATING SYSTEM: DOS 3.30
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Mirabel, Eric P.
REGISTRATION UMBER: 31,211
REFERENCE/DOCKET NUMBER: TN
TELECOMMUNICATION INFORMATION:
               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: PRIOR APPLICATION NUMBER:
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   PPLICANT:
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Local Similarity 100.0%;
hes 8; Conservation
                                                                                                                                                                                                                                                                                   TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE:
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ZIP: 77025
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8; Conserv
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10301 Stella Link Rd.
                                                                                                                                                                                                         100.0%;
nilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                            (713) 664-8914
Leclerc, Claude
Sebo, Peter
                            Ladant, Daniel
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                                                                                                                                                                                                                                                                                                                                                                             (713) 664-2288
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                                                                                                                                                                                                           Score 39; DB 1;
Pred. No. 0.072;
; Mismatches 0
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Pred. No. 0.072;
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US-08-410-360-7
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Best Local Similarity 100.0%;
Matches 8; Conservative 0,
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                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch
COMPUTER: IBM PS-
COMPUTER: 
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NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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LENGTH: 15 amino acids
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 77025
                                                                                                                                                                                                                                                                                                                                                           STATE: Texas
FILING DATE:
                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                              STREET: 10301
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Meyers, Kenneth J.
REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 03495-0109-01000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 07 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1300 1 50
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                10301 Stella Link Rd.
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1300 I Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fung, Michael S.C., Sun, Bill N.C, Sun, Cecily R.Y., Kim, Young W., Yu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ullmann,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tanox Biosystems, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             202-408-4000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                THE GC1q RECEPTOR, HIV-1 gp120 REGION BINDING THERETO, AND RELATED PEPTIDES AND TARGETING ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Recombinant Mutants for Inducing Specific Immune Responses
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                       US/08/410,360
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Pred. No. 0
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                                                                                               TELEFAX: 703-241-2848
TELEX: 248345
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                        FILING DATE: 18-SEP-1991
ATTORNEY/AGENT INFORMATION:
NAME: SVENESON, Leonard R.
REGISTRATION NUMBER: 30,330
REFERENCE/DOCKET NUMBER: 1173
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (713) 664-8914
INFORMATION FOR SEQ ID NO:
                                              SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
   MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/095,332
FILING DATE: 23-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
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NAME: Mirabel, Eric P.
REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TW
TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 301 N. Washington CITY: Falls Church
                                                                                                                                              TELEPHONE:
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les 8; Conserv
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5711947
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                                                                                                                              703-241-2848
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Takahashi, Hidemi
                                linear
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                                                                                                                                                 703-241-1300
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peptide
internal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    METHOD TO INDUCE CYTOTOXIC T LYMPHOCYTES
SPECIFIC FOR A BROAD ARRAY OF HIV-1 ISOLATES USING HYBRID
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                                                                                                                                                                                                                                                            07/760,530
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Pred. No. 0.072;
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                                                                                                                                                                                                                                                                                                                                                            Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 15;
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US-08-707-801A-7
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                                                                                                                                      JS-08-707-801A-7
                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino
                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 7, Application US/08707801A Patent No. 5728814
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                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: DOS 3.30
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/70
FILING DATE: 09/04/1996
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: 08/410,31
FILING DATE: 03/24/1995
ATTORNEY/AGENT INFORMATION:
NAME: Mirabal, Eric P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Fung, I
APPLICANT: Liming
                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
TELEPHONE: (713) 664-2288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Peptide LOCATION: 1..15
OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PS/
OPERATING SYSTEM:
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Local Similarity 100.0%;
Local Similarity 100.0%;
hes 8; Conservative 0
                                                                                                                                                      TYPE: amino acid
TOPOLOGY: linear
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ZIP: 77025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Houston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INDIVIDUAL ISOLATE:
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                                 RAFVTIGK 8
RAFVTICK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Tanox Biosystems, Inc.
10301 Stella Link Rd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fung, Michael S.C., Sun, Bill N.C, Sun, Cecily R.Y., Kim, Young W., Yu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            THE 9C1q RECEPTOR, HIV-1 9p120 REGION BINDING THERETO, AND RELATED PEPTIDES AND TARGETING ANTIBODIES
                                                                  100.0%; Score 39; DB 1; Length 15; 100.0%; Pred. No. 0.072; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "synthetic peptide, sequence = residues 315
to 329 of HIV-1, isolate IIIB, gpl60 envelope
glycoprotein."
                                                                                                                                                                                                                                                                                                                                                                                08/410,360
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US-08-711-175-7
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                                                                                                                                                                                                                                                                                                         Sequence 7, Application US/08711175
Patent No. 5739306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 7, Applic Patent No. 5731428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                       GENERAL INFORMATION:

APPLICANT: Fung, Michael S.C., Sun, Bill N.C, Sun, Cecily R.Y.,
APPLICANT: Kim, Young W., Yu, Liming
TITLE OF INVENTION: THE GC1q RECEPTOR, HIV-1 gp120 REGION BINDING
TITLE OF INVENTION: THERETO, AND RELATED PEPTIDES AND TARGETING
TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 7
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Fung,
APPLICANT: Kim, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: DOS 3.30
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/709,006
FILING DATE: 09-SEP-1996
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/410,360
FILING DATE: 24-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Mirabel, Eric P.
REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TNX95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 77025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: THE GC1q RITITLE OF INVENTION: THERETO, ANTIHODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Houston STATE: Texas
                                                      COUNTRY:
                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                               CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                   ADDRESSEE:
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                                        77025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 RAFVTIGK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
                                                                          Texas
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                                                                                                                 E: Tanox Biosystems, Inc.
10301 Stella Link Rd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10301 Stella Link Rd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fung, Michael S.C., Sun, Bill N.C., Sun, Cecily R.Y., Kim, Young W., Yu, Liming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 39; DB 1; Length 15; ilarity 100.0%; Pred. No. 0.072; Conservative 0; Mismatches 0: Trdolo
                                                          USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Diskette, 3.5 inch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             664-2288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       THE gClq RECEPTOR, HIV-1 gp120 REGION BINDING THERETO, AND RELATED PEPTIDES AND TARGETING
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RESULT 35
US-08-488-252-27
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PRIOR APPLICATION NUMBER: 08/410,360
APPLICATION NUMBER: 08/410,360
FILING DATE: 24-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: MITABEL, ETIC P.
REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TNX95-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (713) 664-2288
TELEPAX: (713) 664-8914
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local (
    PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08\326,676

FILING DATE: 07-Jun-1995

APPLICATION NUMBER: 07\726,605

FILING DATE: 09-July-1991

APPLICATION NUMBER: 07\63,262

FILING DATE: 01-Mar-1991

APPLICATION NUMBER: 07\155,321

FILING DATE: 12-Feb-198

ATTORNEY_AGENT INFORMATION:

NAME: Maria C. H. Lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Chang Yi Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 345 PAKK
CITY: NEW YORK
STATE: NEW YORK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: DOS 3.30
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/488,252
FILING DATE:
CLASSIFICATION: 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
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Local Similarity 100.0%;
nes 8; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 15 amino TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                           FLOPPY DISK
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09-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SYNTHETIC PEPTIDES AND PROCESS
OF USING SAME FOR THE DETECTION OF ANTIBODIES TO
HUMAN IMMUNODEFICIENCY VIRUS (HIV) GP120 ENVELOPE
PROTEIN, DIAGNOSIS OF AIDS AND PRE-AIDS CONDITIONS
AND AS VACCINES
Lin
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; TELEX: 422523 COOPUI
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 15 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-021-879-2
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Best Local Similarity
"arches 8; Conserv?
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                                                                Matches
                                                                                               Query Match
                                                                                                                                                                                                                                                                               COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RELEASE #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/021,879
FILING DATE: 24-FEB-193
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 41189/JPW/AJM
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEPAX: (212) 664-0525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2,
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-758-4800
TELEPHAX: (212) 751-6849
                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
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                            1 RAFVTICK 8
                                                               Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30 Rockefeller Plaza
                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 39; DB 1; Length 15; ilarity 100.0%; Pred. No. 0.072; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Graham P. Allaway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cooper & Dunham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SYNERGISTIC COMPOSITION OF CD4-BASED PROTEIN AND ANTI-HIV-1 ANTIBODY, AND METHODS OF USING SAME
                                                                         100.0%; Score 39; DB 2; 100.0%; Pred. No. 0.072;
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                                                             Mismatches
                                                                                           DB 2; Length 15;
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US-07-760-530-1
                                                                                                               RESULT 38
US-07-950-571A-3
                                                                                                                                                                                                                                                                                                                                             US-07-760-530-1
                                                                                                                                                                                                                                                             Query Match 100.0%; Score 39; DB 2; Best Local Similarity 100.0%; Pred. No. 0.072; Matches 8; Conservative 0; Mismatches
                                                                       Sequence 3, Application US/07950571A Patent No. 5854400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1,
                                                           GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 248345
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30,330
REFERENCE/DOCKET NUMBER: 117:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-241-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: I
FRAGMENT TYPE: I
ORIGINAL SOURCE:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICANT: Chang, Tse Wen, Fung, Michael S.C., Sun, Bill N.C., Sun, Cec APPLICANT: Chang, Nancy T.
TITLE OF INVENTION: Monoclonal Antibodies which Neutralize HIV-1 Infection
                                                                                                                                                                                                                                                                                                                                                            LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
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TITLE OF INVENTION: METHOD TO INDUCE CYTOTOXIC T LYMPHOCYTES

TITLE OF INVENTION: SPECIFIC FOR A BROAD ARRAY OF HIV-1 ISOLATES USING HYBRID

TITLE OF INVENTION: SYNTHETIC PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Falls Church
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 703-241-2848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INDIVIDUAL ISOLATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 POPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH:
                                                         INFORMATION:
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                                                                                                                                                                                                                            1 RAFVTICK 8
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AMINO ACID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19910918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide
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                                                                                                                                                                                                                                                                                                                                                          /note= "synthetic peptide, sequence = residues 315
to 329 of HIV-1, isolate IIIB, gp160 envelope
glycoprotein."
                                                                                                                                                                                                                                                                                                                                                                                                                    /label- peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IIIB
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                                                                                                                                                                                                                                                                                                   DB 2; Length 15;
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                                   Sun, Cecily R.Y
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Best Local Similarity
Workches 8; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 6,
                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: MATSUO
APPLICANT: CHUJO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                       APPLICANT:
            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FO-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Mirabel, Eric P.
REGISTRATION NUMBER: 31,211
REFERENCE/DOCKET NUMBER: TN
                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Hi Density Diskette
                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 10301
CITY: Houston
STATE: Texas
                                                                                                                                     STREET: 1755 S.
CITY: ARLINGTON
STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 199209
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OPERATING SYSTEM:
APPLICATION NUMBER:
                                                                                                                   COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
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                                                                                                          22202
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                                                                                                                                                                                                                                                                                                                                                     MATSUO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  713-664-8914
                                                                                                                                                                                                                                            TASAKA, HIROMICHI
VENTION: ANTI-AIDS SECRETORY RECOMBINANT BCG
VENTION: VACCINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                            HONDA, MITSUO
                                                                                                                                                                                                                                                                                                                           YAMAZAKI,
                                                                                                                                                                                                                                                                                         YAMAKAZI, SHUDO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tanox Biosystems, Inc.
                                                                                                                                                                                               OBLON. SPIVAK, MCCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER: No. 5854400 07/767,533
09/26/1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19920922
                                                                                                                                                                    JEFFERSON DAVIS HIGHWAY, SUITE 400
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                                                                                                                                                                                                                                                                                                                                                        KAZUHIRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DOS, Version 3.30
                                                                                                                                                                                                                                                                                                                         AKIHIRO
US/08/975,699
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RESULT 40
US-08-972-089-6
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Best Local Similarity
Matches 8; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: MATSUO
APPLICANT: CHUJO,
                                                                                                                     COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/972,089
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[NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ent No.
                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: YAMAKAZI, SHUDO
APPLICANT: TASAKA, HIROMICHI
TITLE OF INVENTION: ANTI-AIDS SECRETORY RECOMBINANT BCG
TITLE OF INVENTION: VACCINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide ORIGINAL SOURCE: ORGANISM: HUMAN IMMUI
                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 178462/1994
FILING DATE: 29-UUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                        ADDRESSEE: OBLON. SPIVAK, MCCCLELLAND, MAIER & NEUSTADT, ADDRESSEE: P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400 CITY: ARLINGTON
APPLICATION NUMBER: PCT/JP95/01515 FILING DATE: 31-JUL-1995
                                                                                                        FILING DATE
                                                                                                                                                                                                                                              COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN: HIV-1 (JAPAN)
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
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REGISTRATION NUMBER:
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APPLICATION NUMBER: PCT/JP95/01515
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FILING DATE: 29-MAR-1996
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                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MATSUO,
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                                                 08/975,699
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                NAME: SVENBSON, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                   STREET: P.O. Box 747
CITY: Falls Church
STATE: Virgin:
                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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NAME: OBLON, NORMAN F.
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APPLICATION NUMBER:
FILING DATE: 29-JUL-
TELEFAX:
                                                                                                                                                                             FILING DATE:
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-795-0X PCT
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 703-205-8050
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Ahlers, Jeffrey D.
Pendleton, C. D.
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                                                                                                                                       US 08/060,988
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                                                    1173-434P
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US-08-455-625-11
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Patent No. 5932218
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APPLICANT: Berzof
                                                                                                                                                                                 TELEPHONE: 703-205-800
TELEPAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIDM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                               SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
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                                                                                        OLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REPLICANT: Shirai, Mutsunori
RITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
RITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
RITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
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Local Similarity 100.0%; Pred. No. 0.072;
les 8; Conservative 0; Mismatcher
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STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: 1.15
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                          NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
OTHER INFORMATION: OTHER INFORMATION:
                                  NAME/KEY: Peptide
LOCATION: 1..15
                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: Peptide
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TOPOLOGY: linear
                                                                                                                                                                  ENGTH:
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Ahlers, Jeffrey D.
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/note= "p18IIIB peptide, see Table V"
/label= peptide
/note= "p18-3, see Table V"
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                                                                                                  US-08-455-625-12
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                                   Matches
                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best
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                                                  Best
                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/455,625
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/ACENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                               TELEFAX: 703-205-8050 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                             MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Birch, Ster
STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                             LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
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                                                  Local
                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 22040-0747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                              NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                               IELEPHONE: 703-205-8050
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   1 RAFVTIGK 8
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                                 Similarity
8; Conserv
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VENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
VENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
VENTION: LYMPHOCYTES AGAINST HIV
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                                   Conservative
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                                                                                                                                                                                                peptide
internal
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100.0%; Pred. No. 0.072;
                                100.0%; Score 39; DB 2; Length 15; 100.0%; Pred. No. 0.072; tive 0; Mismatches 0; Indels
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                                                                                                             /label= peptide
/note= "p18-4, see Table V"
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RESULT 45
US-08-455-625-14
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US-08-455-625-13
                                                                                      Sequence 14, Application US/08455625
Patent No. 5932218
                                                                                                                                                                                                                                                                                         Query Match
Best Local (
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                                                                  GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08,
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08,
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: CONTROL OF THE PROPERSON:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM C compatible
COMPUTER: IBM C compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICANT:
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TITLE OF INVENTION: COMPOSIT
TITLE OF INVENTION: LYMPHOCY
                                                                                                                                                                                                                                                                res 8; Conservative 0, Migmath 100.03; Score 39; DB 2; Local Similarity 100.03; Pred. No. 0.072;
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APPLICANT: Ahlers, Jeffrey D
                                                                                                                                                                                                                                                                                                                                                           LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: Peptide LOCATION: 1..15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
TOPOLOGY: linear
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CITY: Falls Church
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5932218
Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. D.
Nara, Peter
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internal
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ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
LYMPHOCYTES AGAINST HIV
                                                                                                                                                                                                                                                                                                                                                    /label= peptide
/note= "p18-5, see Table V"
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                                                                                                                                                                                                                                                                                               DB 2; Length 15;
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RESULT 46
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 703-205-8000
TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                           APPLICANT: Berzofsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
APPLICANT: Pendleton, C. D.
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
                                                          CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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FRAGMENT TYPE:
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ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
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TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
NUMBER OF SEQUENCES: 36
ADDRESSEE: Birch, St
STREET: P.O. Box 747
CITY: Falls Church
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Local Similarity 100.0%;
hes 8; Conservative (
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OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "pl8-6, see Table
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: P.O. Box 7.
CITY: Falls Church
STATE: Virginia
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                                     Stewart, Kolasch & Birch
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SER: 1173-434P
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Pred. No. 0.072;
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                                                                                                                CYTOTOXIC
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; NAME/KEY: Peptide
; LOCATION: 1..15
; OTHER INFORMATION:
; OTHER INFORMATION:
; OTHER INFORMATION:
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Patent No. 5935580
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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NFORMATION FOR SEQ ID NO:
                                       ZIP: 2005-3315

ZOP: 2005-3315

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
LENGTH: 15 amino acid
                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08, FILING DATE: 14-MAY-1993 ATTORNEY AGENT INFORMATION: NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                          RPPLICANT: Ullmann, Agnes
IITLE OF INVENTION: Recombinant Mutants for Inducing
IITLE OF INVENTION: Specific Immune Responses
NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 703-205-8000.
                                                                                                                                                                                       STREET: 1300 I St
CITY: Washington
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CLASSIFICATION: 435
               APPLICATION NUMBER: US/08/395,204 FILING DATE:
                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                           ADDRESSEE:
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E: Dunner
1300 I Street, N.W.
                                                                                                                                                                                                                                                                                                                                        Ladant, Daniel
Leclerc, Claude
Sebo, Peter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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/note= "p18-7, see Table
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US-08-628-687-1
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                                                                                                                                                        TELEFAX: 212-596-9090 INFORMATION FOR SEQ ID NO:
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APPLICANT: Rakowi
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              ANTI-SENSE: NO FRAGMENT TYPE:
                            MOLECUL: P
                                                         TOPOLOGY: li
                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Haley Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: API
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/138,141
FILING DATE: 15-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 03
TELECOMMUNICATION INFORMATION:
                                                                                                                                       SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Rakowicz-Szulczynska, Eva M.
TITLE OF INVENTION: DETECTION AND TREATMENT OF BREAST AND
TITLE OF INVENTION: GYNECOLOGICAL CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 21-APR-1992
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                         TELEPHONE: 212-596-9090
                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/628,687 FILING DATE: 14-JUN-1996
                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
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                                                                                            STRANDEDNESS:
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                                                                                                                          ENGTH:
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                                                                                                        i: 15 amino acids amino acid
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Pred. No. 0.072;
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Best Local S
Matches 8
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: pe
HYPOTHETICAL: NO
FRAGMENT TYPE: in
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CÓMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
                OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                           NAME/KEY: Peptide LOCATION: 1..15 OTHER INFORMATION: OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Margulies, David H.
TITLE OF INVENTION: Potent Peptide for Stimulation of
TITLE OF INVENTION: Cytotoxic T Lymphocytes Specific for the HIV-I Envelope
NUMBER OF SEQUENCES: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tent No.
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                                                                                      NAME/KEY:
LOCATION:
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                                                                                                                                         OTHER INFORMATION:
                                                                                                                                                                                                                                NAME/KEY:
                                                                                                                                                                                                                                                                   STRAIN:
                                                                                                                                                                                                                                                                                    ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30,330
REFERENCE/DOCKET NUMBER: 1173-392P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/07/847,311A FILING DATE: 06-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Birch, Stewart, Kolash & Birch
STREET: 301 N. Washington
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 22046-0747
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8; Conserv
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                                                                                                                                                                                                                                                                               Human Immunodeficiency Virus Type I
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Margulies, David H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pendleton, C.
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Takeshita, Toshiyuki
                                                                                                     Peptide
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N: 435
                                                                                                                                                                                                                                                                                                                                                    peptide
                                                                                                                                                                                                                                                                                                                        internal
/label= peptide
/note= "Highly immunogenic core peptide from
immunodominant region of envelope glycoprotein of
HIV-I strain IIIb; peptide p18-I-10"
                                                                                                                            /label= peptide
/note= "Cytotoxic T lymphocyte immunodominant
/note= "Cytotoxic T lymphocyte immunodominant
peptide of HIV-I envelope glycoprotein from st
IIIB; activatible by protease cleavage to core
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                                                                                                                                                                                                                                    RESULT 51
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; ORGANISM: Human immunodeficiency virus
US-08-986-234-13
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                                         Sequence 28, Application US/08986234
Patent No. 5981706
GENERAL INFORMATION:
APPLICANT: Wallen, et al.
TITLE OF INVENTION: Methods for Synthesizing Heat Shock Protein Complexes
FILE REFERENCE: UNME-0008-1
CURRENT APPLICATION NUMBER: US/08/986,234
CURRENT FILING DATE: 1997-12-05
NUMBER OF SEQ ID NOS: 114
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/08/986,234
CURRENT FILING DATE: 1997-12-05
NUMBER OF SEQ ID NOS: 114
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 8; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Wallen, et al.
TITLE OF INVENTION: Methods for Synthesizing
FILE REFERENCE: UNME-0008-1
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Patent No. 5981706
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                    TYPE: PRT
ORGANISM: Human immunodeficiency virus
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OTHER INFORMATION:
OTHER INFORMATION:
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8; Conserv
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/note= "peptide p18-V-10"
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/note= "peptide
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/note= "peptide
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/note= "peptide p18-I-9"
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                                                                                                                                                                                                                                                                                                                                                   Score 39; DB 2; Length 15; Pred. No. 0.072;
                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Heat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shock Protein Complexes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
                                                                                                                                                                                                                                                                                                                                     Indels
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RESULT 53
US-08-493-071-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6127149
GENERAL INFORMATION:
APPLICANT: Hirai, Yohei
APPLICANT: Koshida, Sho
APPLICANT: Oka, Yumiko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 22, Application US/08492076A
Patent No. 6060064
GENERAL INFORMATION:
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: PC-COMPATIBLE
COMPUTER: PC-COMPATIBLE
COMPUTER: PC-COMPATIBLE
COMPUTER: PC-COMPATIBLE
COMPUTER: US/08/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/493,071
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Price, ROBERT L.
REGISTRATION NUMBER: 22,685
REFERENCE/DOCKET NUMBER: 715-107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 25,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Adams, Sally E.
APPLICANT: Burus, Nigel R.
APPLICANT: Burus, Nigel R.
APPLICANT: Richardson, Simon M.
TITLE OF INVENTION: No. 6060064el Proteinaceous Particles
CURRENT APPLICATION NUMBER: US/08/492,076A
CURRENT FILING DATE: 1995-06-28
EARLIER APPLICATION NUMBER: PCT/GB93/02656
EARLIER APPLICATION NUMBER: PCT/GB93/02656
EARLIER FILING DATE: 1993-12-24
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 15
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: MODIFIED EPIMORPHIN UMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               / Match 100.0%;
Local Similarity 100.0%;
nes 8; Conservative 0
                                                                                                                                                                                                                                                                                         COUNTRY: US
ZIP: 22314
                                                                                                                                                                                                                                                                                                                                  CITY: ALEXANDRIA
STATE: VA
                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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99 CANAL CENTER PLAZA, SUITE 300
                                                                                                                                                                                                                                                                                                                USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/316,436
FILING DATE: 29-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/035,443
FILING DATE: 23-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Zalipsky, Samuel, Woodle, Martin, Martin, Francis;
APPLICANT: Barenholz, Yechekel
                                                                                                                                                                                                                 TELEFAX: (415) 324-09 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 703-684-1124 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
               FEATURE:
NAME/KEY:
                                                                 HYPOTHETICAL: N
ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 7-JUN-1995
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.0%;
hes 8; Conservation
                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 38,563 REFERENCE/DOCKET NUMBER: 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94306
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                                                INDIVIDUAL ISOLATE:
                                                                                                                                                                                                                                                TELEPHONE:
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                                                                                                                LECULE TYPE:
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                                                                                                                                                                                                                                                                                                                      NAME: Mohr,
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30 Cambridge Avenue, Suite 250
                                                                                                                                                                                                                                                    (415)
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                                                                                                                                                 unknown
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                                                Peptide 1, Fig. 13
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US-08-480-332-1

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                                                                                                                                         TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-455-685-7
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Best Local Similarity
Marches 8; Conserve
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                                                                   Matches
                                                                                    Query Match
Best Local (
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INFORMATION FOR SEQ ID NO:
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APPLICANT: Berzofs
APPLICANT: Ahlers,
APPLICANT: Pendlet
                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/0
FILING DATE: 14 MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-MAY-1995
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                                                                                                                                                                                                                                                                                                    TELEPHONE: 617/542-5070
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MUTIIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                   Local Similarity
nes 8; Conserv
                                                                                                                                                                                                                                                                                                                                     NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-4 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
                                                                                                                                                                                                                                                                                   TELEFAX:
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STATE: MA
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                          RAFVTIGK 8
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RAFVTIGK 15
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                                                                                                                                                                                                         15 amino acids
                                                                                                                                                                                                                                                                               617/542-8906
                                                            100.0%; Score 39; DB 3; Length 15; ilarity 100.0%; Pred. No. 0.072; Conservative 0; Mismatches 0; Indels
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Ahlers, Jeffrey D.
Pendleton, C. David
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RESULT 57 US-08-455-685-12

Sequence 12,

Application US/08455685

Patent No. 6214347
GENERAL INFORMATION:
APPLICANT: Berzoff
APPLICANT: Ahlers

Berzofsky, Jay A. Ahlers, Jeffrey D. Pendleton, C. David

APPLICANT:

Nara, Peter

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                                                                                                                                                US-08-455-685-11
                                                                       Query Match
Best Local S
Matches 8
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US-08-455-685-11
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                                                                                                                                                                                                                     TELEX: 200154
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/060
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/748
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148
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MEDIUM TYPE: Diskette
COMPUTER: IBM COMPACTION
OPERATING SYSTEM: Windd
SOFTWARE: FASTSEQ for 1
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                                                                                                                                                                   MOLECULE TYPE:
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TELEPHONE: 617/542-5070
TELEPAX: 617/542-8906
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
                                                                                                                                                                                 LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
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                                                                       Local Similarity
mes 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                 NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Boston
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                                 1 RAFVTICK 8
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                                                                         Conservative
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Pendleton, C. David
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                                                                   100.0%; Score 39; DB 3; Length 15, 100.0%; Pred. No. 0.072; tive 0; Mismatches 0; Indels
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Patent No. 621%...
Parenal INFORMATION:
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COMPUTER: IBM Comparized
OPERATING SYSTEM: Windows95
SOFTWARE: FASESEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
FILING DATE: 31-WAY-1995
PRIOR APPLICATION DATA:
NEDITCATION NUMBER: 08/060,988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/060,988
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                            APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                 APPLICANT:
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REFERENCE/DOCKET NUMBER: 0883
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
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                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                             APPLICANT:
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nes 8; Conserv
                                STREET:
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                                                     ADDRESSEE:
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Boston
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                                    225 Franklin Street
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Ahlers, Jeffrey D.
Pendleton, C. David
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Best Local Similarity
""> hes 8; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: BEATTIE, INGITE A.
REGISTRATION NUMBER: P-42,31
REFERENCE/DOCKET NUMBER: 088
TELECOMMUNICATION INFORMATION:
       COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: Fast-SEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
DATA:
US/08/455,685
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                       COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                              APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compati
                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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FILING DATE: 31-MAY-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/00
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APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
                                                                                                                                                                                    STREET: 225 F
CITY: Boston
STATE: MA
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX:
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Ahlers, Jeffrey D.
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3YSTEM: Windows95
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 31-MAY-1995
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100.0%; Pred. No. 0.072;
tive 0; Mismatches
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RESULT 60
US-08-455-685-15
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Best Local Similarity 100.
Thes 8; Conservative
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FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: 08830/022003
REFERENCE/DOCKET NUMBER: 08830/022003
TELECOMMUNICATION INFORMATION:
TELEPAX: 617/542-8906
TELEPAX: 617/542-8906
                                                                                                     APPLICATION NUMBER: US/08/455, FILING DATE: 31-MAY-1995
PRIOR APPLICATION NUMBER: 08/060,988
PILING DATE: 14-MAY-1993
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
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                                                                                                                                                                                                                                     COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                           APPLICATION NUMBER: 07/847,311 FILING DATE: 06-MAR-1992
                                                                                                                                                                                                                                                                                                                 COUNTRY: US
ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Fish & Att.
STREET: 225 Franklin Street
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Nara, Peter
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Ahlers, Jeffrey D.
                                                                                                                                                                                                                                                        IBM Compatible
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Fish & Richardson P.C.
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                                                                                                                                                                                                                                         Windows95
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US-08-060-988A-7
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                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOTTWARE: FBSETSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION NATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.0%;
Matches 8; Conservative
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                                      TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatibl
OPERATING SYSTEM: Windo
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INFORMATION FOR SEQ ID NO: 15:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: N
TITLE OF INVENTION: T
TITLE OF INVENTION: F
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                               NAME: Beattle, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 0883
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                   FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
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APPLICANT:
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ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 15 amino acids TYPE: amino acid TOPOLOGY: linear
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225 Franklin Street
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Ahlers, Jeffrey D.
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THAT ELICIT
HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-060-988A-7
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US-08-060-988A-11
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Best Local Similarity
Watches 8; Conserve
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                                                                     Query Match 100.0%; Score 39; DB 3; Length 15; Best Local Similarity 100.0%; Pred. No. 0.072; Matches 8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION UMMBER: US/08/060,988A
FILING DATE: 14-WAY-1993
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/847,311
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-WAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
PILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
PILING DATE: 26-JAN-1988
                                                                                                                                                                                                                                                              TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                   TOPOLOGY: 11 MOLECULE TYPE:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 08
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattle, Ingrid A.
REGISTRATION NUMBER: P-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTICETERMINANT PEPTIDES
TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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                                                                                                                                                                                                                                                                                                   TELEFAX:
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ZIP: 02110-2804
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STREET: 22
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                                   RAFVTIGK 8
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Ahlers, Jeffrey D.
Pendleton, C. David
                                                                                                                                                                                       linear
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APPLICANT:

Ahlers, Jeffrey D. Pendleton, C. Davi Nara, Peter

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US-08-060-988A-13
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Sequence 13, Application US/08060988A
PATENT NO. 6294322
GENERAL INFORMATION:
APPLICANT: Berzotsky, Jay A.
APPLICANT: Ahlers, Jeffrey D.
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                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,98
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
ZIP: 02.10-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatib
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEPHONE: 617/542-5070
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APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELDER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
LENGTH: 15 amino acid
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SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
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                                                                                                                                                                                                                                                                        Local Similarity
les 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 15 amino acids TYPE: amino acid TOPOLOGY: linear
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Jeffrey D
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                                                                                                                                                                                                                                                                                                                 Length 15;
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US-08-060-988A-14
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                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/847
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                 APPLICANT: Shirai, MUCSUNOTI
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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TITLE OF INVENTION:
TITLE OF INVENTION:
                                                     CORRESPONDENCE ADDRESS
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                                                                           JUMBER OF SEQUENCES:
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PRIOR APPLICATION DATA:
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OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
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MEDIUM TYPE: Diskette
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                 STREET:
                                       ADDRESSEE:
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REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022001
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ZIP: 02110-2804
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STATE: MA
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Boston
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                 2: Fish & Richardson P.C.
225 Franklin Street
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Ahlers, Jeffrey D.
Pendleton, C. David
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MULTIDETERMINANT PEPTIDES
THAT ELICIT
THAT ELICIT
HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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US-08-060-988A-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                              APPLICANT: NATA, PETER APPLICANT: Shirai, Mutsunori TITLE OF INVENTION: MULTIBETER TITLE OF INVENTION: HELPER T-L TITLE OF INVENTION: NEUTRALIZI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: Windows95
SOFTWARE: FASTSEQ for Windows Version 2.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/148
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,
REFERENCE/DOCKET NUMBER: 0
         OPERATING SYSTEM: Windows95
SOFTWARE: FASESEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
                                                                                                           ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
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ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                       CITY: Boston
                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 / Match 100.0%;
Local Similarity 100.0%;
les 8; Conservative 0
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RAFVTIGK 15
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Ahlers, Jeffrey D.
Pendleton, C. David
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                                                                                     IBM Compatible
SYSTEM: Windows95
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                                                                                                                                                                                                                                                                                                                MULTIDETERMINANT PEPTIDES
THAT ELICIT
HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07/148,692
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Pred. No. 0.
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; LENGTH: 15
; TYPE: PRT
; ORGANISM: HIV
US-09-051-006-8
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                                                                                       RESULT 68
US-09-389-390-1
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; MOLECULE TYPE: peptide
US-08-060-988A-15
Sequence 1, Application US/09389390 Patent No. 6558961 GENERAL INFORMATION: APPLICANT: SARPHIE TITLE OF INVENTION: IMMUNODIAGNOST
                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 39; DB 3; Length 15. Best Local Similarity 100.0%; Pred. No. 0.072;
                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/051,006
CURRENT FILING DATE: 1998-03-30
                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Cheong, Hong-Seok
TITLE OF INVENTION: Liposomes Comprising Peptide Antigens
TITLE OF INVENTION: Derived from X Protein of Hepatitis B virus
FILE REFERENCE: 0136/0E154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 617/542-507
TELEPAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Beattie, Ingrid REGISTRATION NUMBER: F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 07/8 FILING DATE: 06-MAR-1992 APPLICATION NUMBER: 07/7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
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                                                                                                                                                          RAFVTICK 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chang, Jin-Soo
Cho, Sung-Yoo
Hwang, Yu-Kyeong
Choi, Myeong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kim, Tae-Youn
Lee, Ki-Young
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                                                                                                                                                                                                                              Conservative
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     IMMUNODIAGNOSTICS USING PARTICLE DELIVERY METHODS
                                                                                                                                                                                                                                              100.0%; Score 39; DB 3; 100.0%; Pred. No. 0.072;
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US-09-827-688-9
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; TYPE: PRT
; ORGANIZM: Human immunodeficiency virus type 1
US-09-508-552-15
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Sequence 9, Application US/09827688
Patent No. 6821955
GENERAL INFORMATION:
APPLICANT: ORSON, FRANK
APPLICANT: KINSEY, BERNA
APPLICANT: BHOGAL, BERNA
APPLICANT: BHOGAL, BALBIR
TITLE OF INVENTION: MACROAGGREGATED PROTEIN CONJUGATES AS ORAL GENETIC IMMUNIZATION
TITLE OF INVENTION: AGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn V
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn V
SEQ ID NO 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Belyakov, Jay A.
APPLICANT: Belyakov, Igor M.
APPLICANT: Derby, Michael A.
APPLICANT: Kelsall, Brian L.
APPLICANT: Strober, Warren
APPLICANT: THE GOVERNMENT OF THE UNITED STATES OF AMERICA, as
APPLICANT: THE GOVERNMENT OF THE UNITED STATES OF AMERICA, as
TITLE OF INVENTION: MUCOSAL CYTOTOXIC T LYMPHOCYTE RESPONSES
FILE REFERENCE: 368200PCSEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/389,390
CURRENT FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 60/099,261
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: 60/139,045
PRIOR FILING DATE: 1999-06-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/PRIOR FILING DATE: 1998-02-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1997-0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence:synthetic OTHER INFORMATION: construct
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                                                                                                                                                                                                                                                                                           Local Similarity hes 8; Conserv
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les 8; Conserv
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PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60/074,894
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                                                                                                                                                                                                                                                                                             0; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                               DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 4; Length 15;
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PRIOR APPLICATION NUMBER: 60/195,680
PRIOR FILING DATE: 2000-04-07
NUMBER OF SEQ ID NOS: 13
SOPTWARE: PatentIn version 3.0
SEQ ID NO 9
LENGTH: 15
TYPE: PRT
ORGANISM: HIV p18
US-09-827-688-9
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PCT-US92-10378-1
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Best Local Similarity 100.
"--->hes 8; Conservative
                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/US92/1037
FILING DATE: 19921202
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/800,932
FILING DATE: December 2, 1991
PRIOR APPLICATION NUMBER: 07/945865
FILING DATE: September 16, 1992
ATTORNEY/AGENT INFORMATION:
NAME: December 16, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                TELEX: Not Applicable INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: P01949US1/10004014
CURRENT APPLICATION NUMBER: US/09/827,688
CURRENT FILING DATE: 2001-04-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                           NAME: Parker, David L.
REGISTRATION NUMBER: 32,165,
REFERENCE/DOCKET NUMBER: UTI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 512-320-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WORDERAFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: PLATSOUCAS, Chris
APPLICANT: NEHETE, Pramod N.
TITLE OF INVENTION: METHODS
TITLE OF INVENTION: FOR ELIC
STRANDEDNESS: single
TOPOLOGY: linear
OLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                        LENGTH: 15 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Houston
STATE: Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
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                                                                                                                                                               512-474-757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SD
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ARLINGHAUS, Ralph B.
PLATSOUCAS, Chris D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BOARD OF REGENTS, THE UNIVERSITY OF TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arnold,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                METHODS AND COMPOSITIONS
FOR ELICITING IMMUNE OR ANTI-INFECTIVE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 39; DB 4; Length 15; 100.0%; Pred. No. 0.072;
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RESULT 73
PCT-US94-05142-11
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                                                                                                                            Matches
                                                                                                                                             Query Match
Best Local (
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Best Local
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIA Release #1.0,

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                      NAME: SVENBSON, Leonard R.
REGISTRATION INVMER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 14-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                NAME/KEY: Peptide
LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "plBIIIB peptide, see Table V"
                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                             IELEPHONE: 703-205-8050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Birch, St. STREET: P.O. Box 747 CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                           1 RAFVTIGK 8
                                                                                                                       h 100.0%;
Similarity 100.0%;
8; Conservative 0
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                                                           RAFVTIGK 15
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Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                               15 amino acids
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internal
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ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
LYMPHOCYTES AGAINST HIV
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                                                                                                                                        Score 39; DB 5;
Pred. No. 0.072;
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                                                                                                                       Mismatches
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Sequence 11, Application PC/TUS9405142 GENERAL INFORMATION:
APPLICANT:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local Similarity
COUNTS.
ZIP: 22040-074,
ZIP: 22040-074,
COMPUTER READABLE FORM:
COMPUTER: FIODPY disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: FLOPPY disk
COMPACTION OF THE PC-DOS/MS-DOS
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                                                                                                                                                                                                        TITLE OF INVENTION: (
TITLE OF INVENTION: (
TITLE OF INVENTION: (
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ADDRESSEE: Birch, St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                     ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330 REFERENCE/DOCKET NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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STREET: P.O. Box 747
CITY: Falls Church
STATE: Virginia
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                                                                                                                          STATE:
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internal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= peptide
/note= "p18-3, see Table V"
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RESULT 75
PCT-US94-05142-13
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Best Local Similarity
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                                                                                                                                                                                                 ZIF: 22040-0747

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIA RELEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 3
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TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION: TELEPHONE: 703-205-8000
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FRAGMENT TYPE:
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                                                  NAME: Svensson, Leon REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Birch, St
STREET: P.O. Box 747
CITY: Falls Church
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OTHER INFORMATION:
OTHER INFORMATION:
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CLASSIFICATION:
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                               REFERENCE/DOCKET NUMBER: 1173-434P
                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 13-MA
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                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
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TOPOLOGY: linear
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                                                             Leonard R.
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                                                                                                                                                                                                                                                                                                                                                                                     Stewart, Kolasch & Birch
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/note= "p18-4, see Table V"
                                                                                                                  US 08/060,988
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INFORMATION FOR SEQ ID NO. SEQUENCE CHARACTERISTICS

13:

ENGTH:

15 amino acids

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PCT-US94-05142-14
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Best Local
                                                                                         LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acide
                                                                                                                                                                                                                   NAME: Svensson, Leonard R.
REGISTRACION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                  FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
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TITLE OF INVENTION:
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             OTHER INFORMATION:
                                                            NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 22040-0747
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OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-5, see Table V"
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                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
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8; Conserv
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                                                         Peptide
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                                                                                       peptide
internal
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        /label= peptide
/note= "p18-6, see Table V"
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100.0%;

Score 39;

DB 5; Length 15;

US-08-657-392-28 RESULT 78

Sequence 28

28,

B, Application US/08657392 5843634

GENERAL INFOR APPLICANT: APPLICANT:

INFORMATION:

Brate, E.M. Brennan, C.A.

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                                                                                                                                                                                                                                                                PCT-US94-05142-15
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PCT-US94-05142-15
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                                                                                                                                  Matches
                                                                                                                                                                                               Query Match
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Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: lir
MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM CC compatible
COMPUTER: IBM CC compatible
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA;
CURRENT APPLICATION DATA;
COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND COMPAND CO
                                                                                                                                                                                                                                                                                     NAME/KBY: Peptide LOCATION: 1..15
OTHER INFORMATION: OTHER INFORMATION:
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TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: P.O. Box 747
CITY: Falls Church
STRIE: Virgin:
                                                                                                                              Local Similarity
les 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 1
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ELEPHONE:
RAFVTICK 15
                                                             RAFVTICK 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15 amino acids
                                                                                                                              Conservative
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Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide
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                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                              /label= peptide
/note= "p18-7, see Table V"
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ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
LYMPHOCYTES AGAINST HIV
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                                                                                                                                                      Score 39; DB 5;
Pred. No. 0.072;
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                                                                                                                           Mismatches
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                                                                                                                                                                                     Length 15;
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                                                                                                                Gaps
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CITY: NEW YORK

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RESULT 79
US-08-251-472-2
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                                                                                                                                                                                                                                            Sequence 2, Application US/08251472 Patent No. 5871746
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: SOFTEC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/657,392
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/100,708
FILING DATE: July 29, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Wong, Wean Khing
REGISTRATION NUMBER: 5324.US.P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 938-3517
                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: BOUTILI
APPLICANT: FREDER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                      APPLICANT: BOUTILLON, CHRISTOPHE; MARTINON, APPLICANT: FREDERIC; GRAS-MASSE, HELENE; APPLICANT: GOMAND, ELISABETH; SEEGHERAERT, APPLICANT: CHRISTIAN; MAGNE, REMY; TARTAR, APPLICANT: ANDRE; LETY, JEAN-PAUL TITLE OF INVENTION: CYTOTOXIC T LYMPHOCYTE TITLE OF INVENTION: -INDUCING LIPOPEPTIDES NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
CORRESPONDENCE ADDRESS:
ADDRESSEE: BIERMAN & MUSERLIAN
STREET: 600 THIRD AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acid residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Genetically Engineered Enzymes And Their
TITLE OF INVENTION: Conjugates For Diagnostic Assays
NUMBER OF SEQUENCES: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 39; DB 2; Length 16; Local Similarity 100.0%; Pred. No. 0.077; hes 8; Conservative 0. Minnothin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: ABBOTT LABORATORIES
STREET: One Abbott Park Road
CITY: Abbott Park
STATE: Illnois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (708)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 60064-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                           1 RAFVTIGK 8
                                                                                                                                                                                                                                                                                                                                                                     RAFVTICK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Krafft, G.A.
Mandecki, W.
March, S.C.
Russell, J.R.
Yue, V.T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bridon, D.E
Jaffe, K.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28:
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                                                                                   AND USE AS VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/251
PILING DATE: 31-MAY-1994
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: MUSERLIAN, CHARLES A
REGISTRATION NUMBER: 19.683
REFERENCE/DOCKET NUMBER: 102.
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPAN: (212) 661-8000
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 35, App. No. 597655
ZIP: 20005-3315
COMPUTER READABLE FOOM:
MEDIUM TYPE: Floppy Disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS-/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,905
FILING.DATE: 07-JUNE-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/801,818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 10016
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: ASCII
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: peptide FRAGMENT TYPE: internal ORIGINAL SOURCE: ORGANISM: HIV-1
                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Abastado, Jean-Pierre
APPLICANT: Kourilsky, Philippe
TITLE OF INVENTION: An Altered Major Histocompatibility
TITLE OF INVENTION: Complex(MC) Determinant and Methods for Using the
TITLE OF INVENTION: Determinant
NUMBER OF SEQUENCES: 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 16
                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: un....
STRANDEDNESS: un....
STRANDEDNESS: un....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 39; DB 2;
Local Similarity 100.0%; Pred. No. 0.077;
les 8; Conservative 0; Mismatches
                                                                                                                                                                                                                     STATE: D.C.
ZIP: 20005-3315
                                                                                                                                                                                                                                                       STREET: 1300 I St
CITY: Washington
                                                                                                                                                                                                                                                                                              ADDRESSEE: Finneg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: ENV 312-327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RAFVTICK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RAFVTIGK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/08484905
                                                                                                                                                                                                                                                                           1300 I Street, N.W., Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mottez, Estelle
                                                                                                                                                                                                                                                                                                                Finnegan, Henderson, Farabow, Garrett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 16;
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Query Match
Best Local Similarity
Matches 8; Conserve
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US-08-481-985B-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: peptide
US-08-484-905-35
                                                                                                                                         CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/481,985B
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/801,818
FILING DATE: 05-DEC-1991
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/792,473
FILING DATE: 15-NOV-1991
CLASSIFICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION: 435
CLASSIFICATION: 435
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Mottez, Esterne
APPLICANT: Abastado, Jean-Pierre
APPLICANT: Kourilsky, Phillipe
APPLICANT: Kourilsky, Phillipe
APPLICANT: NVENTION: Altered Major Histocompatibility Complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tent No.
                              NAME: Meyers, Kenneth J.
REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 034
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 33,332
REFERENCE/DOCKET NUMBER: 03
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: D.C.
TTP: 20005-3315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Potter, Jane E. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 15-NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     35, AT
NO. 6011146
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dunner
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       202-408-4000
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                                                                                                                                                                                                                                                                                                                        US/08/481,985B
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                                                                     03495.0106-04000
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; LOCATION:
US-09-248-082-2
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Query Match 100.0%; Score 39; DB 3; Length 16; Best Local Similarity 100.0%; Pred. No. 0.077; Matches 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                        TELEFAX: (212) 661-8002
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/25
FILING DATE: 31-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: MUSERLIAN, CHARLES A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                MOLECULE TYPE: peptide FRAGMENT TYPE: internal ORIGINAL SOURCE: ORGANISM: HIV-1
                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 19,683
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 661-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII
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APPLICANT:
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APPLICATION NUMBER: US/09/248;082
FILING DATE:
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: unl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
Local Similarity 100.0%;
nes 8; Conservative 0;
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TYPE: amino acid
TOPOLOGY: linear
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    Application US/09248082
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                                                                                               ENV 312-327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CHRISTIAN; MAGNE, REMY; TARTAR,
                                                                                                                                                                                                        linear
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                                                                                                                                                                                                                          unknown
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Pred. No. 0.077;
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Gaps

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RAFVTIGK 8

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Query Match
Best Local Similarity
Where 8; Conserve
RESULT 84
US-08-992-877-15
; Sequence 15, Ap
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; TOPOLOGY: linear
; MOLECULE TYPE: pep!
US-08-370-476-35
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US-08-370-476-35
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CLASSIFICATION: 435
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/117,575
FILING DATE: 07-SEP-193
APPLICATION NUMBER: US 08/072,787
FILING DATE: 06-JUN-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 202-408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 15-NOV-1991
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                      NAME: Meyers, Kenneth J.
REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 05243.0001-01000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 0 FILING DATE: 05-DEC-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE:
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                                                                                                               1 RAFVTICK 8
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   Application US/08992877
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                                                                                                                                                                                                                                                                       16 amino acids
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Kourilsky, Phillipe
Lone, Yu-Chun
                                                                                                                                               Conservative
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                                                                                                                                                            Score 39; DB 3;
Pred. No. 0.077;
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SOFTWARE: PatentIn V
SEQ ID NO 15
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/
CURRENT FILING DATE: 1997-12-
PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1997-04-17
                                                           ATTORNEY/AGENT INFORMATION:
NAME: Wong, Wean Khing
REGISTRATION NUMBER: 33,561
REFERENCE/DOCKET NUMBER: 532:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 938-3517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Terman, David S
TITLE OF INVENTION: SUPERAN
TITLE OF INVENTION: TRREATN
FILE REFERENCE: superantige
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TYPE: PRT
ORGANISM: Artificial Sequence
NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: SOFEPC
CURRENT APPLICATION DATA:
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TITLE OF INVENTION:
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                                                                                                                                                                       APPLICATION NUMBER:
                                                TELEFAX:
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                                                                                                                                                            CLASSIFICATION:
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Abbott Park
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Brennan, C.A.
Bridon, D.P.
Jaffe, K.D.
Krafft, G.A.
Mandecki, W.
March, S.C.
Russell, J.R.
Yue, V.T.
                                                                                                                                                                                                                                                                                                                        USA
                                                (708)
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Pred. No. 0.077;
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NESULT 87
U$-08-121-054C-3
; Sequence 3, Application US/08121054C
; Patent No. 5637481
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Best Local Similarity
"hes 8; Conserva
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                                                                                                                                                                    Matches
                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                    TELEFAX: (516)273-1717
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/08015770B Patent No. 5683695
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 11/66
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                  NAME: Wilson, M. Lisa
REGISTRATION NUMBER: 34,045
REFERENCE/DOCKET NUMBER: 201
TELECOMMUNICATION INFORMATION:
TELECHONE: (516)273-2828
                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/00
FILING DATE: 10-FEB-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 73
CORRESPONDENCE ADDRESS:

ADDRESSSE: United Biomedical, Inc.
STREET: 25 Davids Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
ORIGINAL SOURCE:
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TITLE OF INVENTION: containing multiple antigenic determinants linked by
TITLE OF INVENTION: flexible domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Shen, De Fen
APPLICANT: Wang, Chang Yi
                                                                                                                                                                   Local Similarity
nes 8; Conserv
                                                                                                                                                                                                                                                                         TYPE: amino acid TOPOLOGY: linear
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STATE: NY
                                                                                                                                                                                                                                                                                                       LENGTH:
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                                                                                                                                  1 RAFVTIGK 8
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                                                                                                                                                             100.0%; Score 39; DB 1; I
100.0%; Pred. No. 0.086;
ative 0; Mismatches 0;
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: FIRM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION UNMBER: US/08/121,054C
FILING DATE: 13-SEP-1993
CLASSIFICATION NUMBER: US 08/013,420
FPICA APPLICATION NUMBER: US 08/013,420
FILING DATE: 01-FEB-1993
ATTORNEY/AGENT INFORMATION:
NAME: Adriano, Sarah B.
REGISTRATION NUMBER: 34,470
REFERENCE/DOCKET NUMBER: 34,470
REFERENCE/DOCKET NUMBER: 30436.18US01
TELEPHONE: 310-445-031
TELEPHONE: 310-445-031
TELEPHONE: 310-445-011
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                                                                                                                                                                                                                                                                                              US-08-488-252-28
                                                                                                                                                                                                  Patent No. 5763100:
GENERAL INFORMATION:
GENERAL INFORMATION:
Chang Yi Wang
TON SYNT
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TITLE OF INVENTION: F
TITLE OF INVENTION: F
TITLE OF INVENTION: 1
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                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                              NUMBER OF SEQUENCES:
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APPLICANT:
APPLICANT:
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CORRESPONDENCE ADDRESS:
                                                       STREET:
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                                        CITY:
                                                                          ADDRESSEE:
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CITY: Los Angeles
STATE: CA
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                                                                                                                                                                                                                                                      28, Applic
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E: NEW YORK
TRY: USA
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                                                     345 PARK AVE.
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Linsley, Peter S.
Bajorath, Jurgen
Fell, Perry
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Gilliland, Lisa K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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                                                                                                          AND AS VACCINES
38
                                                                                                                                       SYNTHETIC PEPTIDES AND PROCESS
OF USING SAME FOR THE DETECTION OF ANTIBODIES TO
HUMAN IMMUNODEFICIENCY VIRUS (HIV) GP120 ENVELOPE
PROTEIN, DIAGNOSIS OF AIDS AND PRE-AIDS CONDITIONS
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Pred. No. 0
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COMPUTER READABLE FORM: MEDIUM TYPE: PLOPPY

PLOPPY DISK

COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WORDPERFECT 5.1

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Best Local Similarity
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TELEX: 421792
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
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FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08\326,676
FILING DATE: 07-Jun-1995
APPLICATION NUMBER: 07\726,605
FILING DATE: 09-July-1991
APPLICATION NUMBER: 07\663,262
FILING DATE: 01-Mar-1991
APPLICATION NUMBER: 07\155,321
FILING DATE: 12-Feb-1988
FILING DATE: 12-Feb-1988
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-758-4800
TELEPAX: (212) 751-6849
TELEX: 421707
                                                                                                                                                 STREET: LILL CITY: LOS Angeles
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                  COUNTRY: USA
ZIP: 90025
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Merchant
                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Expression Vectors Encoding Bispecific
TITLE OF INVENTION: Fusion Proteins and Methods of Producing Biologically
TITLE OF INVENTION: Active Bispecific Fusion Proteins in a Mammalian Cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Maria C. H. Lin
REGISTRATION NUMBER: 29,
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                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: Amino acids
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11150 Santa Monica Blvd., Suite 400
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                                                                                                                                                                                                                                                                                                                                    Fell, Perry
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Linsley, Peter S.
                                                                                                                                                                                                                                                                                                                                                                                                           Ledbetter, Jeffrey A. Gilliland, Lisa K.
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US/08/539,436
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                                    Version #1.30
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PRIOR FILING DATE: 2000-04-13
PRIOR PELICATION NUMBER: 08/539,436
PRIOR PELICATION NUMBER: 08/121,054
PRIOR APPLICATION NUMBER: 08/121,054
PRIOR FILING DATE: 1993-09-13
PRIOR APPLICATION NUMBER: 08/013,420
PRIOR FILING DATE: 1993-02-01
NUMBER OF SEQ ID NOS: 32
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                                                                                Best Local
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SEQ ID NO 3
LENGTH: 20
TYPE: PRT
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                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ledbetter, Jeffrey A.
APPLICANT: Hayden, Martha S.
APPLICANT: Linsley, Peter S.
APPLICANT: Bajorath, Jurgen
APPLICANT: Bajorath, Jurgen
APPLICANT: Gilliland, Lisa K.
TITLE OF INVENTION: EXPRESSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC
TITLE OF INVENTION: FUSION PROTEINS IN A MAMMALIAN CELL
FILE REFERENCE: 30436.18USD2
CURRENT APPLICATION NUMBER: US/09/813,659
CURRENT FILING DATE: 2001-03-21
CURRENT FILING DATE: 2001-03-21
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TELEPAX: 310-445-9031
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
PRILING DATE: 13-SEP-1993
APPLICATION UMBER: US 08/
FILING DATE: 01-FEB-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 30
TELECOMMUNICATION INFORMATION:
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nes 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Adriano, Sarah B.
REGISTRATION NUMBER: 34,470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 05-OCT-1995
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                                       1 RAFVTIGK 8
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                                                                                Similarity
8; Conserv
RAFVTICK 19
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                                                                              100.0%; Score 39; DB 4; ilarity 100.0%; Pred. No. 0.096; Conservative 0; Mismatches
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                                                                                                                   DB 4; Length 20;
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; SOFTWARE: PatentIn Ver.; SEQ ID NO 3; LENGTH: 20; TYPE: PRT; ORGANISM: Homo sapiens US-09-549-067A-3
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Best Local :
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No. 6623940
                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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PRIOR FILING DATE: 1991-06-27
NUMBER OF SEQ ID NOS: 35
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PRIOR APPLICATION NUMBER: 08/539,436
PRIOR FILING DATE: 1995-10-05
PRIOR APPLICATION NUMBER: 08/121,054
PRIOR APPLICATION NUMBER: 08/121,054
PRIOR APPLICATION NUMBER: 08/121,054
PRIOR APPLICATION NUMBER: 08/121,054
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APPLICANT: Gilliland, Lisa K.
TITLE OF INVENTION: EXPRESSION VECTORS ENCODING BISPECIFIC FUSION PROTEINS
TITLE OF INVENTION: AND METHODS OF PRODUCING BIOLOGICALLY ACTIVE BISPECIFIC
TITLE OF INVENTION: FUSION PROTEINS IN A MAMMALIAN CELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 08/013,420 PRIOR FILING DATE: 1993-02-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/549,067A CURRENT FILING DATE: 2000-04-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Linsley, Peter S.
APPLICANT: Bajorath, Jurgen
APPLICANT: Fell, H. Perry
APPLICANT: Gill.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 30436.18USC1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ledbetter, Jeffrey A. APPLICANT: Hayden, Martha S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   quence 4, Application US/08452503A
tent No. 5849475
                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                          IITLE OF INVENTION: Retrovirus-Like Particles Containing
                                                                                                                                                  COUNTRY: Canada
ZIP: M5G 1R7
                                                                                                                                                                                                          STREET: 330 Un
                                                                                                                                                                                                                                                         ADDRESSEE:
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APPLICATION NUMBER: 08/228,208
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 1994-04-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/008,898
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8; Conservative 0
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                                                                                                                                                                                                                            330 University Avenue, 6th Floor
                                                                                                                                                                                                                                                                                                                                                                                              Aliv. Rovinski, ser
                                                                                                                                                                                                                                                                                                                                                             Klein, Michel H
                                 PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                       Sim & McBurney
                                                                                                                                                                                                                                                                                                                                                                                 Shi Xian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AND METHODS OF PRODUCING BIOLOGICAL FUSION PROTEINS IN A MAMMALIAN CELL
                                                                                                                                                                                                                                                                                                                                                                                                                       Benjamin
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                                    TELEFAX: (416) 595-1163 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (416) 595-1163 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                   NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,73
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
FILING DATE: 09-JAN-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0,
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Retrovirus-Like Particles Containing TITLE OF INVENTION: Modified Envelope Glycoproteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/453,745A FILING DATE: 30-MAY-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     330 University Avenue, 6th Floor
21 amino acids
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Cao, Shi Xian
Klein, Michel H
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DN: 435
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                                                                                                                                                                                                                                                                                                               Release #1.0, Version #1.25
                                                                                                                                                                                                          US 08/073,526
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Pred. No. 0.
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RESULT 95
US-08-648-298-18
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US-08-470-419-25
                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/290,105
APPLICATION NUMBER: 08/290,105
FILING DATE: AUGUST 15, 1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-385
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: amino acids
STRANDEDNESS: single
Sequence 18, Application US/08648298 Patent No. 5871990
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                                                                                                                                                                                         Matches
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Patent No. 58663
                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/470,419
FILING DATE: CLASSIFICATION:
PRIOR APPLICATION DATA:
ADDITION NUMBER: 08/200 105
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS IITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
                                                                                                                                                                                     Local Similarity
les 8; Conserv
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STATE: Ontario
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
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STREET: Suite 701, 330 University Avenue
                                                                                                                  14
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linear
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KLEIN, Michel H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAO, Shi-Xian
YAO, Fei-Long
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                                                                                                                                                                                                                         DB 2; Length 21;
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SEQUENCE CHARACTERISTICS:
LENGTH: 21 amin-
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STALL.
STALL.
COUNTRY: Canaua
COUNTRY: Canaua
ZIP: M5G 1R7
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
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APPLICANT: Henrik Clausen
APPLICANT: Eric Paul Bennett
                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                       APPLICANT: PERSSON, ROY
APPLICANT: KLEIN, MICHOL H
TITLE OP INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS RETROVIRUS-LIKE PARTICLES
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/648,298
FILING DATE: 15-JUN-1996
CLASSIFICATION: 435
CLASSIFICATION: 435
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 40
TELECOMMUNICATION INFORMATION
TELEPHONE: 212527700
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                                                                                                                                                                                                                                                                                       APPLICANT:
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                                                                                                                             ADDRESSEE: Sim
STREET: 6TH Fl
CITY: Toronto
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STATE: N.
STATE: N.
10022
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REGISTRATION NUMBER: 38,475
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                                                                                                                                                                                                                                                                                                                                                                                                                                                10 RAFVTICK 17
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8; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 RAFVTICK 8
                                                                                                                                                                                                                                                                                                                                                        Application US/08761828
                                                                                                                                        E: Sim & McBurney
6TH Floor, 330 University Avenue
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805 Third Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 39; nilarity 100.0%; Pred. No. Conservative 0; Mismatch
                                                                                                                                                                                                                                                                   CAO, Shi-Xian
YAO, Fei-Long
                                                                                                                                                                                                                                                                                                       ROVINSKI, Benjamin
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NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION UDATA:
APPLICATION UDATA: US 08/073,526
PILING DATE: 09-JAN-1993
                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/452,520B
FILING DATE: 30-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy
              NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10:
TELECOMMUNICATION INFORMATION:
                                                                                    CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/290,105
FILING DATE: 15-AUG-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Retrovirus-Like Particles Containing TITLE OF INVENTION: Modified Envelope Glycoproteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (416) 595-1155
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                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                      STREET: 330 U
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TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: STEWART, Michael REGISTRATION NUMBER: 24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION:
                                                                                                                                                                                                                                                                                                                                    TRY: Canada
M5G 1R7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Klein, Michel H
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Haynes, Joel
Cao, Shi Xian
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(416)
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                                 1038-446 MIS:as
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Best Local Similarity
Antches 8; Conserva
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                                                                                                                                                                                                      TELEFAX: (416) 595-11 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/290,105
FILING DATE: August 15, 1994
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-385 MIS:jb
                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FOR PORTIBLE
COMPUTER: Floppy disk
COMPUTER: FOR PORTIBLE
COMPUTER: FOR PORTIBLE
COMPUTER: PATENTIA RELEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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                                                                                                                               TOPOLOGY:
                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: Suite CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: Canada
ZIP: M5G 1R7
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                          1 RAFVIIGK 8
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Suite 701, 330 University Avenue
                                                      Conservative
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                                                                                                                            linear
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KLEIN, Michel H
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YAO, Fei-Long
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                                                                    100.0%;
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                                                                    Score 39; DB
Pred. No. 0.1;
                                                      Mismatches
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                                                      Indels
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RESULT 99 US-08-776-949-25 ; Sequence 25, Application US/08776949

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REGISTRATION NUMBER: 24,973
REGISTRATION NUMBER: 1038-673 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEPHONE: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                       RESULT 100
US-08-482-810-25
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TOPOLOGY: linear
TS-08-776-949-25
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                   equence 25, Application No. 6080408
                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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ADDRESSEE: Sim & McBurney
                                                                                                                                                                       TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES MADE NON-
TITLE OF INVENTION: INFECTIOUS BY A PLURALITY OF MUTATIONS
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                        APPLICANT: KLEIN, Michel H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                               AUDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
CITY: Toronto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       / Match 100.0%; Score 39; DB 3; Length 21;
Local Similarity 100.0%; Pred. No. 0.1;
nes 8; Conservative 0; Mismatches 0; Indels
                                                         COUNTRY: Cana
ZIP: M5G 1R7
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ZIP: M5G 1R7
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14 RAFVTIGK 21
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                                                                                      Ontario
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CAO, Shi-Xian
YAO, Fei-Long
                                                                     Canada
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           FILING DATE:

CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/290,105

APPLICATION NUMBER: US 08/290,105

FILING DATE: 15-AUG-194

ATTORNEY/AGENT INFORMATION:

NAME: STEWART, Michael I

REGISTRATION NUMBER: 24,973

REFERENCE/DOCKET NUMBER: 1038-798 MIS:jb

TELECOMMUNICATION INFORMATION:

TELECHONE: (446) 595-1155
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APPLICANT: ROVINS
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                  ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: CAO, Shi-Xian
APPLICANT: YAO, Fei-Long
APPLICANT: PERSSON, ROY
APPLICANT: KLEIN, Michel H
TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES
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NAME: STEMART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US 08/292,967
FILING DATE: 22-AUG-1994
CLASSIFICATION: 424
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APPLICATION NUMBER: US/00
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 6th Floor, 330 University Avenue
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SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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(416) 595-1163
                                                                                                                                                                                                                                                                                                                                                                                                                                           Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ROVINSKI, Benjamin
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                                                                                Query Match
Best Local Similarity
                                                                 Matches
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Patent No. 6342228
GENERAL INFORMATION:
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC FOUNTAINS
COMPUTER: IBM PC FOOMBALDIE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/09/636,805
FILING DATE: 10-Aug-2000
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/027,955
FILING DATE: 23-FEB-1998
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
DEFEDENCE / NOCKET NUMBER: 24,973
DEFEDENCE / NOCKET NUMBER: 24,973
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TYPE: amino acid
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                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 1038-1068 MIS:jb TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS RETROVIRUS-LIKE PARTICLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: ROVINSKI,
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                             1 RAFVTICK 8
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                                                                                                                                                                               LENGTH: 21 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: Canada
ZIP: MSG 1R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Toronto
                                                               Conservative
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YAO, Fei-Long
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                                                          100.0%; Score 39; DB
100.0%; Pred. No. 0.1;
tive 0; Mismatches
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                                                                                             DB 3; Length 21;
                                                            0; Indels
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Query Match
Best Local Similarity
"arches 8; Conserve
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US-09-635-754-25
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APPLICATION NUMBER: 08/292,967
FILING DATE: 22-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-924 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPAX: (416) 595-1155
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 103
US-09-258-128-25
                                                                                                                                                                                                       Sequence 25, Application US/09635754
Patent No. 6518030
GENERAL INFORMATION:
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Patent No. 645132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/258,128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: amino acid
STRANDEDNESS: single
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TITLE OF INVENTION: INFECTIOUS BY A PLURALITY OF MUTATIONS
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & MCBUrney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                            APPLICANT: ROVINSKI, Benjamii
CAO, Shi-Xian
YAO, Fei-Long
PERSSON, ROY
KLEIN, Michel H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
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                                                              NUMBER OF SEQUENCES:
                                                                                               TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
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KLEIN, Michel H
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                                                                              RETROVIRUS-LIKE PARTICLES
                                                                                                                                                                                        Benjamin
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Pred. No. 0.1;
Mismatches
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RESULT 105
US-08-680-525-25
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMMBER: US/08/680,525
FILING DATE:
CLASSIFICATION EATA:
APPLICATION NUMBER: US 08/292,967
FILING DATE: 22-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 25, Application US/08680525 Patent No. 6544527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: CAO, Shi-Xian
APPLICANT: YAO, Fei-Long
APPLICANT: PERSSON, ROY
APPLICANT: KLEIN, Michel H
TITLE OF INVENTION: RETROVIRUS-LIKE PARTICLES MADE NON-
TITLE OF INVENTION: INFECTIOUS BY A PLURALITY OF MUTATIONS
                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 26
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FILING DATE: 10-Aug-2000
CLASSIFICATION CUMRIOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/027,955
FILING DATE: 23-FEB-198
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-1065 MIS:jb
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                           COUNTRY: Canada
ZIP: M5G 1R7
                                                                                                                                                                                                                                                                                                 CITY: Toronto
STATE: Ontario
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ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Toronto
STATE: Ontario
                                                                                                                                                                                                                                                                                                                                               B: Sim & McBurney
Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ROVINSKI, Benjamin
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RESULT 106
US-09-636-223-25
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Patent No. 6544732
GENERAL INFORMATION:
APPLICANT: ROVINSKI, Benjamin
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Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 39; DB 4; Best Local Similarity 100.0%; Pred. No. 0.1;
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INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                     INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
ENGTH: 21 amino acids
TYPE: amino acid
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NAME: STEWART, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                    PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 09/027,955

PILING DATE: 23-FEB-1998

ATTORNEY/AGENT INFORMATION:

NAME: STEWART, Michael I

REGISTRATION NUMBER: 24,973

REFERENCE/DOCKET NUMBER: 1038-1064 MIS:jb

TELECOMMUNICATION INFORMATION:

TELEPHONE: (416) 595-1153

TELEFAX: (416) 595-1163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 21 amino acids
TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       POPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                             SEQUENCE DESCRIPTION: SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: ANTIGENICALLY-MARKED NON-INFECTIOUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14 RAFVTICK 21
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                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/636,223 FILING DATE: 29-Dec-2000 CLASSIFICATION: <Unknown>
                                                                              STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: Canada
ZIP: M5G 1R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Toronto
STATE: Ontario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Sim & McBurney
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (416)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAO, Shi-Xian
YAO, Fei-Long
PERSSON, Roy
KLEIN, Michel H
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100.0%; Score 39;
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DB 4;
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Length 21;
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Best Local Similarity 100.0%; Pred. No. 0.1; Matches 8; Conservative 0; Mismatches

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Indels

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Gaps

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APPLICANT:

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                                                                        US-08-783-818-13
                                                                                             RESULT 108
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US-08-125-012-13
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            Sequence 13, Application US/08783818
Patent No. 5817637
GENERAL INFORMATION:
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Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 8; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 215-568-3429
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: APP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/125, FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/029,336
FILING DATE: 11-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/008,342
FILING DATE: 26-JAN-1993
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APPLICANT: Coney, Leslie R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Genetic Immunization TUMBER OF SEQUENCES: 34
                                                                                                                                                                                                                                                                                                                                              LENGTH: 22 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5593972ris
STREET: One Liberty Place 46th Floor
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                     15 RAFVTIGK 22
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5. 5593972
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Williams, William V.
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RESULT 109
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Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.0%; Score 39; DB 2; Best Local Similarity 100.0%; Pred. No. 0.11; Matches 8; Conservative 0; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 215-568-3429
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                           NUMBER OF SEQUENCES: 3
                                                                                                                                                                APPLICANT: Williams, APPLICANT: Wang, Bin TITLE OF INVENTION: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: APTELECOMMUNICATION INFORMATION: TELEPHONE: 215-568-3100
COMPUTER READABLE FORM
                                                                                                                                                                                                                       APPLICANT:
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PRIOR APPLICATION NUMBER: 08/008,342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/1
FILING DATE: 21-SEP-1993
APPLICATION NUMBER: 08/0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: Genetic Immunization
                                 COUNTRY:
                                                                      ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5830876ris
STREET: One Liberty Place 46th Floor
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/0 FILING DATE: 26-JAN-1993 CLASSIFICATION: 514
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CLASSIFICATION: 514
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                   19103
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T: One Liberty Place 46th Floor
Philadelphia
                                                    Pennsylvania
                                                                                                                                                                                                                                                                             Application US/08453349
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                                     USA
                                                                                                                                                                                               Weiner, David B.
Williams, William V.
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                                                                                                                                                                  Genetic Immunization
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Best Local Similarity 100.0%;
Matches 8; Conservative 0;
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APPLICANT: ZOLLA-1
ADDITCANT: ZORNY
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/345,321
ATTORNEY/AGENT INFORMATION:
NAME: Browdy, Roger L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: ZO.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: APP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
                                                                                                           FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                       STREET: 417 COUNTY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/029,336
FILING DATE: March 11, 1993
APPLICATION NUMBER: 08/008,342
FILING DATE: January 26, 1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: GORNY, MIYOBIAY K.
TITLE OF INVENTION: HETERCHYBRIDOMAS PRODUCING HUMAN
TITLE OF INVENTION: MONOCLONAL ANTIBODIES TO HIV-1
NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOPTWARE: PatentIn Release #1.0, Version #1.25 mb-MD/JAF CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                     COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                               FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                              419 Seventh Street, N.W.,
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                     ZOLLA-PAZNER1B
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Pred. No. 0.11;
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RESULT 111
US-08-979-385B-11
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 248633
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/979,385B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF INVENTION: Compositions and Methods for Delivery of NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
                                             APPLICATION NUMBER: FILING DATE: 15-JUL-PRIOR APPLICATION DATA:
                                                                                                                                                                APPLICATION NUMBER: PCT/L
FILING DATE: 26-JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/11
FILING DATE: 21-SEP-1993
                                                                                                                                                                                                                                                 FILING DATE: 26-NOV-1997
CLASSIFICATION: 514
PRIOR APPLICATION UNMBER: US 08/495,684
FILING DATE: 28-SEP-1995
PRIOR APPLICATION UNMBER: US 08/495,684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                               PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
            APPLICATION NUMBER: 08/029,336 FILING DATE: 11-MAR-1993
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OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: One Liberty Place 46th Floor CITY: Philadelphia
                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RAFVTIGK 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Weiner, David B.
Williams, William V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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internal
                                                               15-JUL-1993
                                                                                                               21-SEP-1993
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to 303 to 324 of gp120 from the IIIB isolate."
                                                                                                                                   08/124,962
                                                                                                                                                                                                                                      PCT/US94/00899
                                                                             08/093,235
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Pred. No.
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   Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Barth, Richard
REGISTRATION NUMBER: 28,180
REFERENCE/DOCKET NUMBER: 9501
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 972-1400
TELEPAX: (212) 370-1622
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                      NFORMATION FOR SEQ ID NO: 1:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acide
                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        quence 1, Appli
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                     MOLECULE TYPE:
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TELEFAX: \-\
236268
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Neurath, A. Robert, Debnath, Asim K.,
APPLICANT: Jiang, Shibo
TITLE OF INVENTION: Proteins and Peptides Modified By
TITLE OF INVENTION: Aromatic Acid Anhydride Compounds
NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: UP TELECOMMUNICATION INFORMATION:
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                                                                                                                LENGTH: 22 amino acids
TYPE: amino acid
                                                                                                  STRANDEDNESS: single stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Match 100.0%; Score 39; DB 2 Local Similarity 100.0%; Pred. No. 0.11; es 8; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
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600 Third Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Diskette, 3+ inch, 0.72 mb storage
IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             215-568-3100
                                                                  cDNA to genomic RNA
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 Score 39;
Pred. No.
DB 2;
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              Length 22;
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Best Local Similarity
Matches 8; Conserv
                                                                                                           Sequence
                                                                                         Sequence 5, Application US/09070291
Patent No. 6171596
                                                           GENERAL INFORMATION:
APPLICANT: Earl,
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INFORMATION FOR SEQ ID NO:
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   TITLE OF INVENTION:
                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 36,516
REFERENCE/DOCKET NUMBER: NI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-235-650
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PRIOR APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENELIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Fuller, Michael L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Oligomeric HIV-1 Envelope Glycoproteins NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Broder,
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STATE: ...
92660
PRAD/
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                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 22 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0 FILING DATE: 03-MAR-1997
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                                                                                                                                                                                                                 1 RAFVTICK 8
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T: 620 Newport Center Drive 16th Floor
Newport Beach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8; Conservative
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                 Doms, Robert W. Moss, Bernard
                                            Broder, Christopher C.
                                                              Earl, Patricia L
                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                   linear
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Oligomeric HIV-1 Envelope Glycoproteins
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NIH079.001A
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                                                                                                                                                                                                                                                           Score 39; DB 3
Pred. No. 0.11;
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NUMBER OF SEQUENCES: 1

ADDRESSEE:

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Query Match
Best Local Similarity
""" has 8; Conserve
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Best Local Similarity
"---hes 8; Conserve
                                                                                                                   ORGANISM: Homo sapiens
FEATURE:
FEATURE:
NAME/KEY: MISC_FEATURE
OTHER INFORMATION: HIVIIIB 9p120
US-09-217-306B-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ANTI-SENSE: NO; FRAGMENT TYPE: US-09-070-291-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-217-306B-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 115
                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/217,306B
CURRENT FILING DATE: 1998-12-21
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn version 3.1
SEQ ID NO 22
LENGTH: 22
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22, Application US/09217306B Patent No. 6465220
                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                          APPLICANT: Hassan, Helle
APPLICANT: Clausen, Henrik
APPLICANT: Bennett, Eric P.
TITLE OF INVENTION: Glycosylation Using GalNAc-T4 Transferase
FILE REFERENCE: 8850*1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: peptide HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 619-235-0176
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                          TYPE: PRT
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LENGTH: 22 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Vensko, Nancy Ways
REGISTRATION NUMBER: 36,298
REFERENCE/DOCKET NUMBER: NII
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-235-8550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Vensko, Nancy Ways
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STATE: C.
STATE: P2660
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CLASSIFICATION:
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1 RAFVTIGK 8
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                                                   Conservative
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                                                 100.0%; Score 39; DB 4
100.0%; Pred. No. 0.11;
tive 0; Mismatches
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100.0%; Pred. No. 0.11;
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                                                                                 DB 4;
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                                                   0; Indels
                                                                                 Length 22;
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                                                                                                                                                                                                                                                                                                                   US-08-097-751-1
                                                                                                                                                                                                                                                                                                                                                        RESULT 117
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                                                                                                                                                                                                                                      Sequence 1, Application US/08097751 Patent No. 5527666
                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 646898;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR AFFECTION NUMBER: 08/125, ULZ
APPLICATION NUMBER: 08/029, 336
APPLICATION NUMBER: 08/029, 336
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/008, 342
                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 215-568-3429
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: APOL-0013
TELECOMMUNICATION INFORMATION:
                                            APPLICANT:
                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
        APPLICANT:
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STREET: One Liberty Place 46th Floor
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
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Mammano, Fabrizio
Panozzo, Marina
Dettin, Monica
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Williams, William V.
                                                                                                                 DeRossi, Anita
Pasti, Marcella
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Best Local (
                                                                                                                                                                                                       APPLICANT: Mastico, Robert Allan
APPLICANT: Stockley, Peter George
APPLICANT: Talbot, Simon John
TITLE OF INVENTION: Antigen-Presen
TITLE OF INVENTION: Fusion MS2-Coal
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE Rosenman & Colin
                                                                                    COUNTRY: U.S.A.
ZIP: 10022-2585
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5", 1.44Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (212) 302-89
NFORMATION FOR SEQ ID NO:
                        COMPUTER: IBM PS2-486
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
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LENGTH: 24 amino acids
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NAME: Costigan, James V.
REGISTRATION NUMBER: 25,669
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STREET: LILL
CITY: New York
STATE: New York
TISA
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TELECOMMUNICATION INFORMATION:
TELEPHNE: (212) 302-8989
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SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette,
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FILING DATE:
                                                                                                                                                                           STREET: 575 Mar
CITY: New York
               APPLICATION NUMBER:
                                                                                                                                                               STATE:
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Local Similarity 100.0%;
les 8; Conservative 0
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TYPE: amino acid
                                                                                                                                                                                    ADDRESSEE: Rosenman & COLL.
STREET: 575 Madison Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 199307
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                                                                                                                                                                                                                                                                                                                                              INFORMATION:
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1185 Avenue of the Americas
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Chieco-Bianchi,
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08/11/93
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                                                                                                                                                                                                                                                            Fusion MS2-Coat Protein
                                                                                                                                                                                                                                                                          Antigen-Presenting Capsid with
                                                                                                                                                                                                                                                                                                         Peter George
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METHOD FOR T
            US/08/090,148
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Pred. No. 0.11;
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US-08-257-528B-99
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                                                                              TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 99
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 AMINO ACIDS
TYPE: AMINO ACID
TOPOLOGY: NOT RELEVANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/257,528B
FILING DATE: 09-UN-1994
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART MICHAEL I.
                                                                                                                                                                                                            NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9101550.3
FILING DATE: 01/24/91
APPLICATION NUMBER: PCT/GB92/00124
FILING DATE: 01/22/92
ATTORNEY/AGENT INFORMATION:
NAME: NIFORMATION:
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REGISTRATION NUMBER: 27,882
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Toronto
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Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (212) 940-8636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Charles D.Y.
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ER: 27,582
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Pred. No.
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100.0%;

Score 39; Pred. No.

DB 1; 0.11;

Length 24;

Tandem Synthetic HIV-1 Peptides

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                                                                                                        US-08-463-966A-99
                                                                                                                      RESULT 121
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                                                                  Sequence 99, Application US/08463966A Patent No. 5795955
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                                                      GENERAL INFORMATION:
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TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
PILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA: 08/073,378
APPLICATION NUMBER: 08/073,378
PILING DATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY/ACENT MICRAFICATION:
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APPLICANT: SIA, C
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                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
TYPE: amino acid
STRANDEDNESS: single
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                   APPLICANT:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460,602A
FILING DATE: 02-JUN-1995
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APPLICANT: KLBIN, Michel H.
TITLE OF INVENTION: Tandem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                          / Match 100.0%;
Local Similarity 100.0%;
nes 8; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-450 MIS:jb
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ZIP: M5G 1R7
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                                                                                                                                                                         14 RAFVTIGK 21
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SIA, Charles D.Y
CHONG, Pele
KLEIN, Michel H.
                                                                                                                                                                                                                                                                                                                                  linear
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                                  Charles D.Y.
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Pred. No. 0.11;
Mismatches
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US-08-465-217A-99
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION MUMBER: US/08/463,966A
FILING DATE: 05-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/257,528
                                                                                                                                                                                                                                                                                                                                    Patent No. 5800822
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                           Sequence 99, Application US/08465217A
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INFORMATION FOR SEQ ID NO:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPANIAN
OFTHARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
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REFERENCE/DOCKET NUMBER: 1038-487 MIS:jb
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 08/257,526
FILING DATE: 09-UN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/073,378
                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
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                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                 APPLICANT: SIA,
                                                                                                                                                                                                                                                                FITLE OF INVENTION:
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CITY: Toronto
STATE: Ontario
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Local Similarity 100.0%; Pred. No. 0.11;
hes 8; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                 STATE: Ontario
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                                                                                                                          OUNTRY:
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ite 701, 330 University Avenue
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                     Version #1.25
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                         CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY/ACENT INFORMATION:
NAME: STEWART, MICHAEL I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,329A
FILING DATE: 05-JUN-1995
                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                          CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/2:
FILING DATE: 09-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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APPLICANT: KLBIN, Michel H.
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LENGTH: 24 amino acids
TYPE: amino acid
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NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-486 MIS:jb
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
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APPLICATION NUMBER: 08/073,378
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REFERENCE/DOCKET NUMBER: 1038-449 MIS:jb
                  REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                 ZIP: M5G 1R7
                                                                                                                                                                                                                                                                                                                                                                                        STREET: Suite
CITY: Toronto
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TOPOLOGY: linear
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Suite 701, 330 University Avenue
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nilarity 100.0%;
Conservative 0
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, 595-1163
, NO: 99:
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               24,973
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Pred. No. 0.11;
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US-08-493-235-24
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Best Local Similarity
Matches 8; Conserva
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                    Query Match
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, '
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/493,235
FILING DATE: 20)June)195
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Kaiser, Annemarie
REGISTRATION NUMBER: 37,649
REFERENCE/DOCKET NUMBER: METRICS.
                                                                   HYPOTHETICAL: NATI-SENSE: NO FRAGMENT TYPE:
                                                                                                                                                                                                                 TELEFAX: 619-235-0176
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                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Horal, Peter
TITLE OF INVENTION: PEPTIDES FOR USE IN VACCINATION AND
TITLE OF INVENTION: INDUCTION OF NEUTRALIZING ANTIBODIES AGAINST HUMAN
TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS
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                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                  LENGTH: 24 amino acids
TYPE: amino acid
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                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U:
ZIP: 92660
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TYPE: amino acid
STRANDEDNESS: single
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5840313
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Horal, Peter
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                                                                                                  NO NO
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                                                                      internal
                                                                                                                                                   single
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100.0%; Score 39; DB 2; 100.0%; Pred. No. 0.11;
                                                                                                                                                                                                                                                                                                                                                                                                                   Release #1.0, Version #1.25
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                 Length 24
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Best Local Similarity
Matches 8; Conserva

Conservative

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Mismatches

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Indels

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Gaps

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                                                                                                                                                                      RESULT 126
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                                                                                                               Sequence 160,
Patent No. 58
                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/462,507A
FILING DATE: 05-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION OPERATION: 424
PRIOR APPLICATION UMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
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                                                                                            GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                    / Match 100.0%;
Local Similarity 100.0%;
mes 8; Conservative 0
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CITY: Toronto
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                                                                                                                                                                                                                           14 RAFVTICK 21
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                                                                                                             60, Application US/08146028
5891640
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (416)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SIA, Charles D.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sim & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tandem Synthetic HIV-1 Peptides
PROCESS FOR THE DETERMINATION OF PERFIDES
CORRESPONDING TO IMMUNICACICALLY ANTIBODIES OR BIOTINYLATED IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES,
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                                                                                                                                                                                                                                                                                                                       Score 39; DB 2
Pred. No. 0.11;
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                                                                                                                                                                                                                                                                                                                                           DB 2;
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                                                                                                                                                                                        COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.25

SOFTWARE: PATENTIN Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/467,881A

FILING DATE: 06-JUN-1995

CLASSIFICATION: 424

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/257,528

FILING DATE: 09-JUN-1994

CLASSIFICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: 08/073,378

FILING DATE: 09-JUN-1993

CLASSIFICATION: 424

ATTORNEY/AGENT INFORMATION:

NAME: STEMART MYCHAFL. I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                        TELEFAX: (416) 595-1163 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: SIA, C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: peptide
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COMPUTER READABLE FORM:
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NUMBER OF SEQUENCES:
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APPLICANT: KLEIN, Michel H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                       SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE:
                                                                                                                                        REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-488 MIS:jb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 100.0%; Score 39; DB 2; Length 24; Local Similarity 100.0%; Pred. No. 0.11;
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                                                                                                                                                                              NAME: STEWART, MICHAEL I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: M5G 1R7
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                ENGTH:
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: 24 amino acids amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              453
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STRANDEDNESS:

TOPOLOGY: linear

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RESULT 129
US-08-480-332-2
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US-08-723-425A-160
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                                                                                    Sequence 2, Application US/08480332 Patent No. 6180134
                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                          Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                            Query Match
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Best Local Similarity 100.0%;
Matches 8; Conservative 0
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                                  GENERAL INFORMATION:
APPLICANT: Zalips
APPLICANT: Barenh
                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 140
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/723,425A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 22201
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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ADDRESSEE: NIXON & VANDERHYE, P.C.
STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS: 811
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: SADOFF, B.J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH:
CANT: Zalipsky, Samuel; Woodle, Martin; Martin, Francis; CANT: Barenholz, Yechezkel
OF INVENTION: Enhanced Circulation Effector Composition
OF INVENTION: Method
                                                                                                                                                                                         15
                                                                                                                                                                                                                    1 RAFVTIGK 8
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                                                                                                                                                                                                                                                    100.0%; Score 39; ilarity 100.0%; Pred. No. Conservative 0; Mismatch
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                             single
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            Enhanced Circulation Effector Composition and
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Pred. No. 0.11;
                                                                                                                                                                                                                                                      Mismatches
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US-09-112-206-160
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                                                                                                                                                                                                                                                        Sequence 160, Application US/09112206
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0860
TELEPAX: (415) 324-0960
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PILING DATE: 29-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/035
PILING DATE: 23-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Mohr, Judy M.
REFERENCE/DOCKET NUMBER: 38,563
REFERENCE/DOCKET NUMBER: 5356
                                                               NUMBER OF SEQUENCES: 4:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                    TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                  TITLE OF INVENTION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
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ADDRESSEE: Dehlinger & Associates
                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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nes 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 24 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION:
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                                                                                                                                                                                                                      INFORMATION:
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ARE: PatentIn Release #1.0, Version #1.25 (EPO)
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                                                                                                                             PROCESS FOR THE DETERMINATION OF PEPTIDES
CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR (
IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED
PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES, I
                                                                                                                  PROCESS FOR PREPARING
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Pred. No. 0.11;
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                                                                                                                  THEM AND COMPOSITIONS CONTAINING THEM
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; LENGTH: 24 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-09-112-206-160
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                                                                                                                              RESULT 132
US-09-790-497A-160
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CURRENT FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: 09/576,824
PRIOR PILING DATE: 2000-05-23
PRIOR PILING DATE: 2000-05-23
PRIOR FILING DATE: 1996-09-30
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR PILING DATE: 1993-11-22
PRIOR APPLICATION NUMBER: PCT/EP93/00517
PRIOR PILING DATE: 1993-03-08
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 14
LENGTH: 24
TYPE: PRT
              Sequence 160, Application US/09790497A
PAtent No. 6649735
GENERAL INFORMATION:
APPLICANT: De Leys, Robert
TITLE OF INVENTION: PROCESS FOR THE D
TITLE OF INVENTION: TO IMMUNOLOGICAL
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: EP 92400598.6 PRIOR FILING DATE: 1992-03-06 NUMBER OF SEQ ID NOS: 600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TITLE OF INVENTION: TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR USE IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 2752-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: De Leys,
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                                                                                                                                                                                                                                                                                             Local Similarity
nes 8; Conserv
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Similarity 100.0%;
8; Conservative 0
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Robert
PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR USE IN
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Pred. No. 0.11;
                                                                                                                                                                                                                                                                                                 Mismatches
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US-09-576-824A-160
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SEQ ID NO 160
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PRIOR APPLICATION NUMBER: 09/576,824
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 08/723,425
PRIOR FILING DATE: 1996-09-30
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR FILING DATE: 1993-11-22
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TITLE OF INVENTION:
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Best Local
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                                                                                                   Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 160,
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PRIOR FILING DATE: 1996-09-30
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR FILING DATE: 1993-11-22
PRIOR APPLICATION NUMBER: PCT/EP93/00517
PRIOR FILING DATE: 1993-03-08
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NUMBER OF SEQ ID NOS: 600
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/576,824A CURRENT FILING DATE: 2000-05-23
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PRIOR APPLICATION NUMBER: EP 92400598.6
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                                                                                                                                                                                 LENGTH: 24
TYPE: PRT
ORGANISM: Human immunodeficiency virus
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nes 8; Conserv
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RAFVTIGK 22
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                                                                                Conservative
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TO IMMUNOLOGICALLY IMPORTANT ENTITOPES AND THEIR USE IN
A PROCESS FOR DETERMINATION OF ANTIBODIES OF
BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY
BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY
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                                                                                                   100.0%; Score 39; 100.0%; Pred. No.
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                                                                                                     0.11;
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                                                                                                                     DB 4; Length 24;
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CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 744,281
FILING DATE: 13 August 1992
ATTORNEY/AGENT INFORMATION:
NAME: Paul T. Clark
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SENERAL INFORMATION:
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APPLICATION NUMBER: US/08/146,028
FILING DATE: 22-NOV-1993
INFORMATION FOR SEQ ID NO: 160:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                       SOFTWARE: Wordberfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NIMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
COMPUTER: IBM PS/2 Model 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: REPLIGEN CORPORATION
APPLICANT: THE ROCKEFELLER UNIVERSITY
TITLE OF INVENTION: MULTIPLE ANTIGEN PEPTIDES FOR USE AS HIV
TITLE OF INVENTION: VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: ITITLE OF SEQUENCES:
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/680,497
FILLING DATE: 06-OCT-2000
DEFICE APPLICATION OFFICE OFFICE APPLICATION OFFICE                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
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    REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US92/06688 FILING DATE: 19920811
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ZIP: 02110-2804
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CITY: Boston
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TYPE: amino acid
STRANDEDNESS: single
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CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR
IN A PROCESS FOR DETERMINATION OF ANTIFODIES OR BIOTINYLATED
PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES,
PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
30,162
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OS (Version 3.30)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Applicati
GENERAL INFORMATION:
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Best Local
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/94ff
PILING DATE: cc-
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INFORMATION FOR SEQ ID NO:
                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
                                                                                                                                                                                                                                                                                                       TELEPHONE: 512-320-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10:
FILLING DATE: 19921202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 07/945865
FILING DATE: September 16, 1992
ATTORNEY/AGENT INFORMATION:
      TOPOLOGY: li
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                        TELEPHONE: 514-J.,
TELEPHONE: 512-474-7577
TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL TOTAL
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COUNTRY: US
ZIP: 77210
                                                                                                                                                                                                                                                                                                                                                                     NAME: Parker, David L.
REGISTRATION NUMBER: 32.
REFERENCE/DOCKET NUMBER:
                                                                        STRANDEDNESS:
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STATE: Texas
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                                                                                                    AMINO ACID
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100.0%; Pred. No.
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ER: UTFC305PCT
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Chris D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 24;
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                                                                                                                                                                                                      US-08-266-448-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/07950571A Patent No. 5854400
                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                              Sequence 1, Application US/08266448 Patent No. 5876724
                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: NO. 58544
PILING DATE: 09/26/1991
ATTORNEY/AGENT INFORMATION:
NAME: Mirabel, Eric P.
REGISTRATION NUMBER: 31,211
                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 713-664-8914
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                   APPLICANT: GIRARD, Marc
TITLE OF INVENTION: INDUC
TITLE OF INVENTION: INFEC
TITLE OF INVENTION: AND P
TITLE OF INVENTION: THE G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Chang, Tse Wen, Fung, Michael S.C., Sun, Bill N.C., Sun, Cec: APPLICANT: Chang, Nancy T.
TITLE OF INVENTION: Monoclonal Antibodies which Neutralize HIV-1 Infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: TN
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-664-2288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Hi Density Diskette
COMPUTER: IBM ps/2
OPERATING SYSTEM: DOS, Version 3.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: DOS, Version 3.30 SOFTWARE: Wordperfect 5.1 CURRENT APPLICATION DATA:
                                    NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 100.0%;
Local Similarity 100.0%;
tes 8; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US
FILING DATE: 19920922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 77025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                    8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tanox Biosystems
FINNEGAN, HENDERSON, FARABOW, GARRETT &
                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                  INDUCTION OF PROTECTION AGAINST VIRAL
INFECTION BY SYNERCY BETWEEN VIRUS ENVELOPE GLYCOPROTEIN
AND PEPTIDES CORRESPONDING TO NEUTRALIZATION EPITOPES OF
THE GLYCOPROTEIN
23
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Pred. No. 0.11;
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Pred. No. 0.12;
                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                   Sequence 13, Application US/08485324 Patent No. 6043093
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Best Local
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/145,664
PILING DATE: 04-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/782,241
FILING DATE: 28-OCT-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/672,647
PRILING DATE: 18-MAR-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/494,749
PRILING DATE: 19-MAR-1990
PILING DATE: 19-MAR-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (202) 408-44
INFORMATION FOR SEQ ID NO:
                                                 ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: not relevant MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
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MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                               APPLICANT: Wohlstadter, Jacob
TITLE OF INVENTION: SELECTION METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 1300 I St
CITY: Washington
                                                                                                                                                                ADDRESSEE: Curtis, Morris,
ADDRESSEE: c/o Barry Evans
STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/01
FILING DATE: 28-JUN-1994
             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Meyers, Kenneth J. REGISTRATION NUMBER: 25,146
                                                                                                           COUNTRY:
                                                                                                                                  STATE:
                                                                                                                                                  CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                        15 RAFVTIGK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RAFVTICK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8; Conservative
                                                                                                                              New York
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                                                                                                              USA
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PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  not relevant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 39; DB 2; 100.0%; Pred. No. 0.12;
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                                                                                                                                                                                                          Morris, & Safford
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Patent No. 6043093
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                                                                                                                                              APPLICATION NUMBER: US/08/485,324
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: US. 07/852,412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (212) 840-3333
TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                          NAME: EVADS, BATTY
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: SE NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
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FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370
TELECOMMUNICATION INFORMATION:
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                                        TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 25 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
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: New York
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                                        (212) 840-3333
(212) 840-0712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wohlstadter, Jacob
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                                    840-0712
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                                                                                                370132-2000
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US-08-447-506-13
                                                                   Matches
                                                                                  Query Match
Best Local
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Patent No. 60
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                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Evans, Barry
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-200
                                                                                                                                                                                                TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Wohlst
                                                                                                                                                                                                                                                                    TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 23-MAY-CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris, & Safford
ADDRESSEE: c/o Barry Evans
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                       Local Similarity 100.0%;
nes 8; Conservative
                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                LENGTH: 25 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 530 Fifth Avenue CITY: New York
STATE: New York
                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: sir
15 RAFVTICK 22
                                1 RAFVTICK 8
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                                                                                                                                                ss: single
linear
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JENTION: SELECTION METHODS
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                                                                            Score 39; DB 3
Pred. No. 0.12;
                                                                Mismatches
                                                                                              Length 25
                                                            0;
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US-08-447-506-31 RESULT 142

; Sequence 31, Application US/08447506
; Patent No. 6066499
; GENERAL INFORMATION:

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US-08-235-437-13
; Sequence 13, Appli
; Patent No. 6087177
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FILING DATE: 23-MAY-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: EVANS, BARTY
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0,

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/235,437
                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                      tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOPTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                              APPLICANT: Wohlstadter, Jacob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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Local Similarity 100.0%;
hes 8; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 25 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                CITY: New York
STATE: New York
                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: New York
STATE: New York
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ADDRESSEE: C/o Barry Evans
                                                                                                                                                                                                                        ADDRESSEE:
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                                                                                                                                               COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                          Application US/08235437
                                                                                                                                                                                                       530 Fifth Avenue
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                                                                                                                                                   USA
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                                    PatentIn Release #1.0, Version #1.25
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 US/08/235,437
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US-08-235-437-31
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US-08-235-437-31
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Best Local Similarity 100.0%;
Query Match
Best Local Similarity
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                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Evans, Barry
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                            TELEFAX: (212) 840-0712 INFORMATION FOR SEQ ID NO:
                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acide
                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 29-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: EVANS, BATTY
REGISTRENCE DOCKET NUMBER: 22,802
REFERENCE DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 07/852,412 FILING DATE: 16-MAR-1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                            TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 25 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U
ZIP: 10036
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CITY: New York
STATE: New Yor
                                                                                      STRANDEDNESS:
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                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Curtis, Morris, & Safford
ADDRESSEE: c/o Barry Evans
STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15 RAFVTIGK 22
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                                                                                                       1: 25 amino acids amino acid
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                                                                     linear
                                                                                    single
100.0%;
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Score 39; DB 3
Pred. No. 0.12;
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Pred. No.
                DB 3;
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                                                                                                               Sequence 31, Application US/08447515
Patent No. 6162640
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/447,515
FILING DATE: 23-MAY 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY AGENT INFORMATION:
NAME: EVANS, BATTY
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
TITLE OF INVENTION: SELECTION METHODS

NUMBER OF SEQUENCES: 31

CORRESPONDENCE ADDRESS:

ADDRESSEE: Curtis, Morris, & Safford

ADDRESSEE: C/o Barry Evans

STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 55 CITY: New York
CITY: New York
TTE: New York
TTSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 37
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                   APPLICANT: Wohlstadter, Jacob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
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NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Wohlstadter, Jacob
                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 25 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                    Local Similarity
nes 8; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Pred. No. 0.12;
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Query Match
Best Local Similarity
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                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 69
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 31
LENGTH: 25
                                                                   Matches
                                                                                 Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 31, Application US/09593870A Patent No. 6548643
                                                                                                                                                                                                                                                   APPLICANT: Apostolopoulos, Vasso
APPLICANT: Pietersz, Geoff Allan
TITLE OF INVENTION: Antigen Carbohydrate Compounds and Their
TITLE OF INVENTION: Use in Immunotherapy
FILE REFERENCE: 2368-McKenzie
CURRENT APPLICATION NUMBER: US/09/593,870A
CURRENT FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: 09/223,043
PRIOR FILING DATE: 1998-12-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,437
FILING DATE: 29-APR-1994
APPLICATION NUMBER: US 07/852,412
FILING DATE: 16-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: EVANS, BAITY
REGISTRATION NUMBER: 22,802
REFERENCE/DOCKET NUMBER: 370132-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (212) 840-07
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: McKenzie, Ian F.C.
                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 10036
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 25 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 23-MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FELEPHONE:
12 RAFVTICK 19
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                                 1 RAFVTIGK 8
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                                                                 Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        : (212) 840-3333
(212) 840-0712
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-MAY-1995
                                                                                 100.0%;
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                                                                                Score 39; DB 4
Pred. No. 0.12;
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Pred. No.
                                                                   Mismatches
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                                                                                                DB 4; Length 25;
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                                                                 0,
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                                                                   Indels
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                                                                 Gaps
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APPLICANT:

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RESULT 148
US-08-455-625-17
                                                                                                                     RESULT 149
US-08-455-625-23
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                                                                                   Sequence 23, Application US/08455625 Patent No. 5932218
                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                            Query Match
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Patent No. 5932218
                GENERAL INFORMATION:
APPLICANT: Berzof
APPLICANT: Ahlers
APPLICANT: Pendle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/455,625
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Berzof
APPLICANT: Ahlers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 703-205-8050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                               NAME/KEY: Peptide LOCATION: 1..15 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 WOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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CITY: Falls Church
STATE: Vireinia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                   B9.7%;
Local Similarity B7.5%;
Local Similarity B7.5%;
Local Similarity B7.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH:
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Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. D.
Nara, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 amino acids
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Ahlers, Jeffrey D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide
internal
                                                                                                                                                                                                                                                                                                                                             /label= peptide
/note= "p18-9, see Table V"
                                                                                                                                                                                                                                                                              Score 35; DB 2;
Pred. No. 0.51;
                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                              Length 15;
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Matches 7; Conserv
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                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08,
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: COMITITLE OF INVENTION: ELIC
TITLE OF INVENTION: LYMINUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                      APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                         tent No.
                                                    APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                     APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: Peptide
LOCATION: 1.15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-15, see Table V"
                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Birch, St
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
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Boston
MA
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                                                                                                                                                                                                                                                                                                            Application US/08455685
                                    225 Franklin Street
                                                                                                                                                                                                          Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. David
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VENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
VENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
VENTION: LYMPHOCYTES AGAINST HIV
                                                                                                                                                                                            Nara, Peter
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Pred. No. 0.51;
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COMPUTER READABLE FORM:

02110-2804

COUNTRY:

MEDIUM TYPE:

Diskette

IBM Compatible SYSTEM: Windows95

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Best Local Similarity
7; Conserve
                                                    COUNTRY: US
ZIP: 02110-2804
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
""""EME Compatible
"""EME: Windows95
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/060,988
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-198
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  equence 23,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tent No.
            SOPTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/455,685
                                                                                                                                                                                                                                                                                              APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/01
FILING DATE: 31-MAY-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                        TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                          CITY: Boston
STATE: MA
                                                                                                                                                                                                                          ADDRESSEE: Fish & Richardson P.C. STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-4 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM CO.
OPERATING SYSTEM:
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6214347
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Ahlers, Jeffrey D.
Pendleton, C. David
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31-MAY-1995
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US-08-060-988A-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDET
TITLE OF INVENTION: THAT ELI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: p-44
REFERENCE/DOCKET NUMBER:
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                            APPLICATION NUMBER:
                                                                         APPLICATION NUMBER: 07/8
FILING DATE: 06-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
TYPE: a
                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                             COUNTRY: US
ZIP: 02110-2804
                                                                                                                                                                                                                                                                                          CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION:
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                                                                                                                                                                                                                                                                                                          225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Berzofsky, Jay A.
                                                                                                                                                                                                                                                                                                                          Fish & Richardson P.C.
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                                            29-AUG-1991
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THAT ELICIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jeffrey D.
                                                                                                                                                                                                                                                                                                                                                       HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          89.7%;
                              07/148,692
                                                         07/751,998
                                                                                        07/847,311
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Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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Best Local Similarity
                                                                                                                                                                                                                      COMPUTER READALE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: PastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION UNUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/184,692
FILING DATE: 26-JAN-198
ATTORNEY/AGENT INFORMATION:
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                                                                      TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tent No.
              SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
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LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 617/542-5070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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                                                                                                                                                                   REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022001
                                                                                                                                                                                          NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
ZIP: 02110-2804
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ropology:
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                                                                                                             617/542-8906
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Ahlers, Jeffrey D.
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87.5%;
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Query Match
Best Local Similarity
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                                                               Matches
                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                 TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                         NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330 REGISTRATION NUMBER: 11. REFERENCE/DOCKET NUMBER: 11. TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                            MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Falls Church
                                                      B9.7%;
Local Similarity 87.5%;
hes 7; Conservative
                                                                                                                                            NAME/KEY: Peptide
LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-9, see Table V"
                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 13-MA'CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                             ENGTH:
                              1 RAFVTIGK 8
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RVFVTIGK 15
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                                                                                                                                                                                                                                               peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCT/US94/05142
                                                                                                                                                                                                                                                                                                                                                                                                  1173-434P
                                                               0; Mismatches
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Pred. No.

    Mismatches

                                                                                Score 35; DB 5
Pred. No. 0.51;
                                                                                               DB 5;
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                                                                                               Length 15;
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RESULT 155
PCT-US94-05142-23
; Sequence 23, Application PC/TUS9405142

ENERAL INFORMATION:

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RESULT 156
US-08-279-906A-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 87.5
                                                                                                                                                                                                                              Sequence 19, Application US/
Patent No. 5618922
GEMERAL INFORMATION:
APPLICANT: Ohno, Tsuneya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/US
FILING DATE: 13-MAY-1994
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
          STREET: 65.
STREET: 65.
CITY: Chicago
CITY: Illinois
TISA
                                                                                                                           APPLICANT: Terada, Masaki
APPLICANT: Yoneda, Yukio
TITLE OF INVENTION: NM03 Antibody Materials and Methods
NUMBER OF SEQUENCES: 27
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: Peptide
LOCATION: 1.15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-15, see Table V"
                                                                                          ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Borun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid TOPOLOGY: linear
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90909
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                                                                        6300 Sears Tower,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC TLYMPHOCYTES AGAINST HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US 08/060,988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCT/US94/05142
                                                                        233 S. Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 35; DB 5; Length 15; Pred. No. 0.51; 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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RESULT 157
US-08-704-170-51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 51,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                        CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: US 08/029
APPLICATION NUMBER: US 08/029
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920
TELECOMMUNICATION INFORMATION:
TELECHONE: (213) 977-1001
    TELEFAX: (213) 977-10
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Takenama, Glenn
APPLICANT: Ehresmann, Glenn
APPLICANT: Ehresmann, Glenn
APPLICANT: THERAPEUTIC STRATEGIES FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite 500
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: NO. 5618922and, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 32028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Douvas, Angeline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 87.5%; Pred. No. 1987 7; Conservative of the cons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 19 amino acids TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A. ZIP: 90012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 RIFVIIGK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RAFVIIGK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/08704170
: (213)
(213) 977-1003
~~~ TD NO: 51:
  IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
  US 08/029,850
  US/08/279,906A
   1920-331
   32028
   DB 1; Length 19;
  0,
  Gaps
```

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PCT-US95-03236-25
; Sequence 25, Application PC/TUS9503236
; GENERAL INFORMATION:
; APPLICANT: University of Southern (
   당
  S
  밁
  ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-704-170-51
   RESULT 158
PCT-US94-02631-51
  RESULT 159
  PCT-US94-02631-51
  Sequence 51, Application PC/TUS9402631 GENERAL INFORMATION:
   Best Local Similarity
Matches 7; Conserv
  Query Match
   Matches
   Query Match
   APPLICATION NUMBER: PCT/US94/02631
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
  TELEFAX: (213) 977-10
INFORMATION FOR SEQ ID NO:
  SEQUENCE CHARACTERISTICS:
   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
  MOLECULE TYPE: peptide
  APPLICANT:
  TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
  NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS: Robbins,
  APPLICANT:
   IITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
   NAME: Spitals, John P. REGISTRATION NUMBER: 29, REFERENCE/DOCKET NUMBER:
   CITY: Los Angeles
STATE: California
   Local Similarity 100 nes 7; Conservative
   TYPE: amino acid
TOPOLOGY: linear
  STREET:
   COUNTRY:
   LENGTH:
   LENGTH: 8 amino acids
  90012
  2 RAFVTIG 8
  1 RAFVTIG 7
   RAFVTIG 7
   amino acid
   8 amino acids
  E: Robbins, Berliner & Carson
201 North Figueroa Street, Suite 500
 University of Southern California
  Douvas, Angeline
Takehana, Yoshi
   U.S.A.
  Ehresmann,
  Conservative
   Floppy disk
  977-1003
   87.2%; Score 34; DB 5;
100.0%; Pred. No. 4.1e+C
tive 0; Mismatches
   100.0%;
   87.2%; Score 34; 100.0%; Pred. No.
  Glenn
   29,215
   0
   1920-331
   Mismatches
  DB 1;
   4.1e+05
   4.1e+05
   0; Indels
  Length 8
  Length 8;
  Indels
  ,
,
   0
  Gaps
   Gaps
  0
   0,
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RESULT 160
  US-08-704-170-38
   , MOLECULE TYPE: peptide
PCT-US95-03236-25
   Sequence 38, App. -- 80, 5707626
   Query Match
Best Local S
  Matches
  GENERAL INFORMATION:
   TELEFAX: (619) 535-89
INFORMATION FOR SEQ ID NO:
   APPLICANT:
APPLICANT:
   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 11-MAR-1993
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
  NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
   APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
   SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
  ATTORNEY/AGENT INFORMATION:
NAME: Imbra, Richard J.
REGISTRATION NUMBER: 37,643
   NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
ATREET: 4370 La Jolla Village Drive, Suite 700
   REFERENCE/DOCKET NUMBER: FP
TELECOMMUNICATION INFORMATION:
   TITLE OF INVENTION:
  CITY: Los Angeles
STATE: California
  STREET:
  TYPE: amino acid TOPOLOGY: linear
   CLASSIFICATION: 424
   FILING DATE:
  APPLICATION NUMBER:
   APPLICATION NUMBER: PCT/US95/03236 FILING DATE: 13-MAR-1995 CLASSIFICATION:
  CITY: San Diego
STATE: California
  COUNTRY:
  TELEPHONE:
  COUNTRY:
  1 RAFVTIG 7
  l Similarity
7; Conserve
   90012
   92122
   RAFVTIG 8
  Application US/08704170
  201 No. 5707626th Figueroa Street, Suite 500
  U.S.A.
  (619) 535-90
(619) 535-8949
  Conservative
  USA
   Takehana, Yoshi
   Douvas, Angeline
  535-9001
   100.0%;
   87.2%; Score 34; DB 5;
100.0%; Pred. No. 4.1e+
   Infection
   Methods to Diagnose and Treat HIV-1
                   US 08/029,850
  US/08/704,170
  ,.
  FP-SI 1394
  Mismatches
   Length 8;
  Indels
  0
  Gaps
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ATTORNEY/AGENT INFORMATION:

REGISTRATION NUMBER:

REFERENCE/DOCKET NUMBER: 1920-331

29,215

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S
   PCT-US94-02631-38
  ; MOLECULE TYPE: peptide US-08-704-170-38
  PCT-US94-02631-38
   Sequence 38, Applicat GENERAL INFORMATION:
  Matches
                             Query Match 87.2%;
Best Local Similarity 100.0%;
Matches 7; Conservative (
   Query Match
   TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO:
   TELEFAX: (213) 977-100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
   FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
APPLICATION NUMBER: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
  SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02631
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
   NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
   MOLECULE TYPE; peptide
  STREET: ZVI ...
STREET: ZVI ...
CITY: Los Angeles
CTATE: California
TT S.A.
  NUMBER OF SEQUENCES: 1:
   APPLICANT:
  TELECOMMUNICATION INFORMATION:
   y ration 87.2%; Score 34;
Local Similarity 100.0%; Pred. No.
hes 7; Conservative 0; Mismarch
  ITLE OF INVENTION:
   TOPOLOGY:
   TYPE: amino acid
  FELEPHONE:
   TOPOLOGY:
   ENGTH:
  DDRESSEE:
   1 RAFVTIG 7
   1 RAFVTIG 7
   amino acid
  Application PC/TUS9402631
   201 North Figueroa Street, Suite 500
   9 amino acids
   Ehresmann, Glenn
VENTION: THERAPEUTIC STRATEGIES FOR
VENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
   Takehana, Yoshi
   Douvas, Angeline
  Robbins,
  linear
  (213)
   977-1003
  977-1001
   US .08/029,850
  Berliner & Carson
   38:
                                  0; Mismatches
  Score 34; pred. No.
   Mismatches
   DB 5;
  4.1e+05;
   Length 9;
   Length 9;
   Indels
                                    Indels
   0,
                                  0,
                                  Gaps
   Gaps
   0
                                  0
```

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   Query Match
Best Local Similarity
"hes 7; Conserv?
   RESULT 162
US-08-704-170-71
  밁
  , MOLECULE TYPE: peptide US-08-704-170-71
   CT-US94-02631-71
   Patent No.
  Sequence 71, Application PC/TUS9402631 GENERAL INFORMATION:
   INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 10 amino acids
  GENERAL INFORMATION:
  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
         CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 North Figueroa Street, Suite 500
CITY: Los Angeles
   REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
   NUMBER OF SEQUENCES:
   TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
   APPLICANT:
   APPLICANT:
  APPLICANT:
  APPLICANT:
  STREET: 201 NO. 50
CITY: Los Angeles
STATE: California
   TYPE: amino acids
  TOPOLOGY:
  NAME: Spitals, John P. REGISTRATION NUMBER: 29,215
  CLASSIFICATION:
  APPLICATION NUMBER:
   ADDRESSEE:
   FILING DATE:
   COUNTRY:
  1 RAFVTIG 7
   RAFVIIG
  Application US/08704170
California
  201 No. 5707626th Figueroa Street, Suite 500
   Douvas, Angeline
Takehana, Yoshi
  Conservative
   U.S.A.
  Ehresmann, Glenn
  (213) 977-1003
   Douvas, Angeline
   Robbins, Berliner & Carson
   UMBER: US 08/029,850
11-MAR-1993
  THERAPEUTIC STRATEGIES FOR IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
   100.0%;
   87.2%; Score 34; DB 1; 100.0%; Pred. No. 0.56;
   121
  US/08/704,170
  0
   .920-331
  Mismatches
  <u>,</u>
  Length 10;
  <u>;</u>
  Gaps
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   RESULT 164
US-08-704-170-73
   PCT-US94-02631-71
  Query Match
Best Local Similarity 100.
  Sequence 73, App. No. 570762
   CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/029
APPLICATION NUMBER: US 08/029
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920
TELEPHONE: (213) 977-1001
TELEPHONE: (213) 977-1003
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920-331
  GENERAL INFORMATION:
   TELEFAX: (213) 977-100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite 500
CITY: Los Angeles
STATE: California
   APPLICANT:
   MOLECULE TYPE: peptide
   SOPTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
   COMPUTER READABLE FORM:
  NUMBER OF SEQUENCES:
  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
  APPLICATION NUMBER:
  TLE OF INVENTION: THERAPEUTIC STRATEGIES FOR INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
  FILING DATE:
   APPLICATION NUMBER: PCT/US94/02631
   FILING DATE:
   ZIP: 90012
   COUNTRY:
   TOPOLOGY:
  COUNTRY: U.S.A. ZIP: 90012
  LENGTH: 10 amino acids
  1 RAFVTIG 7
  amino acid
   Application US/08704170
   Ehresmann,
   U.S.A.
   Takehana, Yoshi
  Douvas, Angeline
   87.2%; Score 34; DB 5; Length 10; 100.0%; Pred. No. 0.56;
   Glenn
  US/08/704,170
   US 08/029,850
   0; Mismatches
   1920-331
   <u>.</u>.
   0
   Gaps
   0
```

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; LENGTH: 11 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-704-170-73
   Query Match
Best Local Similarity
7; Conserve
밁
                             S
  밁
   US-08-704-170-74
   US-08-704-170-74
   RESULT 165
  Query Match
Best Local Similarity
Matches 7; Conserv
  PRIOR APPLICATION DATA:

PRIOR APPLICATION NUMBER: US 08/029

FILING DATE: 11-MAR-1993

ATTORNEY/AGENT INFORMATION:

NAME: Spitals, John P.

REGISTRATION NUMBER: 29,215

REFERENCE/DOCKET NUMBER: 1920

TELECOMMUNICATION INFORMATION:

TELEPHONE: (213) 977-1001
  TELEFAX: (213) 977-100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
  GENERAL INFORMATION:
  TELEFAX: (213) 977-10
INFORMATION FOR SEQ ID NO:
  MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,170
   MOLECULE TYPE: peptide
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
  TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS NUMBER OF SEQUENCES: 121
   SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
  CORRESPONDENCE ADDRESS:
  ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite
CITY: Los Angeles
STATE: California
   TOPOLOGY:
   TYPE: amino acid
   ZIP: 90012
  FILING DATE:
   TELEPHONE:
   CLASSIFICATION:
  COUNTRY:
  LENGTH:
                                 1 RAFVTIG 7
  1 RAFVTIG 7
  5 RAFVTIG 11
   1, Application US/08704170
5707626
  11 amino acids
   Conservative
   (213) 977-1003
   Takehana, Yoshi
Ehresmann, Glenn
   Douvas, Angeline
   Conservative
   (213) 977-1001
  977-1003
  87.2%;
  87.2%; Score 34; DB 1; 100.0%; Pred. No. 0.62;
   US 08/029,850
  73:
  0
  1920-331
   0; Mismatches
  Score 34; DB 1;
Pred. No. 0.62;
  Mismatches
  0;
   0; Indels
  Length 11,
  Length 11;
  0
   0
   Gaps
   Gaps
  0
   0,
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밁
   PCT-US94-02631-73
  PCT-US94-02631-74
   RESULT 167
   Sequence 74, Applicat GENERAL INFORMATION:
   Matches
  Query Match
  equence 73, Applicat
GENERAL INFORMATION:
  TELEFAX: (213) 977-10
INFORMATION FOR SEQ ID NO:
   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: PCT/US94/02631
APPLICATION NUMBER: PCT/US94/02631
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
  APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
  SEQUENCE CHARACTERISTICS:
   REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
  APPLICATION NUMBER: US 0 FILING DATE: 11-MAR-1993 ATTORNEY/AGENT INFORMATION:
   CLASSIFICATION:
PRIOR APPLICATION DATA:
   APPLICANT:
   STREET: ZUL....
STREET: Los Angeles
CITY: Los Angeles
CTATE: California
   CORRESPONDENCE ADDRESS:
   MOLECULE TYPE: peptide
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   NUMBER OF SEQUENCES:
   TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR INFECTIONS
  NUMBER OF SEQUENCES:
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   COUNTRY: U
ZIP: 90012
   NAME: Spitals, John P. REGISTRATION NUMBER: 29,215
   FILING DATE:
  ADDRESSEE:
  ZIP: 90012
   COUNTRY:
  IOPOLOGY:
   DDRESSEE:
   ENGTH:
  1 RAFVTIG 7
  Los Angeles
California
  amino acid
   Application PC/TUS9402631
  Application PC/TUS9402631
   11 amino acids
   E: Robbins, Berliner & Carson
201 North Figueroa Street, Suite 500
  E: Robbins, Berliner & Carson
201 North Figueroa Street, Suite 500
  Ehresmann,
  U.S.A.
   U.S.A.
   Takehana,
   Conservative
   Douvas, Angeline
  linear
  3) 5;
977-1003
977: 73:
   87.2%; Score 34; DB 5; Length 11; 100.0%; Pred. No. 0.62;
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APPLICATION NUMBER: PCT/US95/03236
FILING DATE: 13-MAR-1995
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 37,643
TREGISTRATION NUMBER: 37,643
TORNET NOTOTETT NUMBER: 37,643
   Sequence 29, Applicat GENERAL INFORMATION:
  TELEFAX: (619) 535-8949 INFORMATION FOR SEQ ID NO:
  INFORMATION FOR SEQ ID NO:
  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
APPLICATION NUMBER: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
  NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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TOPOLOGY: linear
MOLECULE TYPE: peptide
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  RAFVTIG 11
   Application PC/TUS9503236
   4370 La Jolla Village Drive, Suite 700
  (213)
   University of Southern California
VENTION: Methods to Diagnose and Treat HIV-1
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  USA
   PatentIn Release #1.0, Version #1.25
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  (619)
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100.0%;
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87.2%;
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RESULT 170
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APPLICANT: Douvas,
APPLICANT: Takehana
APPLICANT: Ehresman
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   TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
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NAME: Imbra, Richard J.
REGISTRATION NUMBER: 37,643
REFERENCE/DOCKET NUMBER: FP
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Local Similarity 100.0%;
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CITY: San Diego
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  ADDRESSEE:
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: California
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   Application US/08704170
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4370 La Jolla Village Drive, Suite 700
                               E: Robbins, Berliner & Carson
201 No. 5707626th Figueroa Street, Suite 500
  USA
  University of Southern California
   Takehana, Yoshi
Ehresmann, Glenn
  Douvas, Angeline
   Floppy disk
  Infection
   Methods to Diagnose and Treat HIV-1
  Score 34;
; Pred. No.
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  Mismatches
  DB 5;
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SOFTWARE: PatentIn Release #1.0, Version #1.25
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APPLICATION NUMBER: 29,215
REGISTRATION NUMBER: 29,215
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
   PRIOR APPLICATION DATA:
APPLICATION NUMBER:
   APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSIT
TITLE OF INVENTION: ELYMPHOCY
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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TELEPHONE: (213) 977-1001
  APPLICANT:
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   STREET: P.O. Box 7
CITY: Falls Church
STATE: Virginia
   APPLICATION NUMBER:
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  ZIP: 22040-0747
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  FILING DATE:
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   1 RAFVTIG 7
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   Virginia
  Application US/08455625
   Berzofsky, Jay A. Ahlers, Jeffrey D. Pendleton, C. D.
   Conservative
   Nara, Peter
   Birch, Stewart, Kolasch & Birch
O. Box 747
                                       14-MAY-1993
  87.2%;
   LYMPHOCYTES AGAINST HIV
  COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
   US 08/060,988
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ATTORNEY/AGENT INFORMATION:
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   SOFTWARE: PatentIn Release #1.0, Version #1.25
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   CORRESPONDENCE ADDRESS
   APPLICANT: Pendleton, C. D.
APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: BLICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
   TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
MOLECULE TYPE:
  STREET: P.O. Box 747
CITY: Falls Church
STATE: Virnini
  Local Similarity nes 7; Conserv
  TELEPHONE: //-
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703-205-8050
TO NO:
   REFERENCE/DOCKET NUMBER: 1173-434P
               TYPE: amino acid
TOPOLOGY: linear
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  COUNTRY:
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   NAME/KEY:
  ENGTH:
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OGY: linear
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5932218
   AFVTIGK 15
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  87.2%; Score 34; DB 2; Length 15; ilarity 100.0%; Pred. No. 0.84; Conservative 0; Mismatches
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  US 08/060,988
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Query Match
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APPLICATION NUMBER: US/08/455,625
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REFERENCE/DOCKET NUMBER: 11/3-434P
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LOCATION:
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  ADDRESSEE: Birch, St. STREET: P.O. Box 747 CITY: Falls Church STATE: Virginia
   TELEPHONE:
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  1 RAFVTICK 8
   H: 15 amino acids amino acid
  Application US/08455625
  Shirai, Muteumori
Shirai, Muteumori
VENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
VENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
VENTION: LYMPHOCYTES AGAINST HIV
  Ahlers, Jeffrey D. Pendleton, C. D.
   Conservative
  Nara, Peter
  Berzofsky, Jay A.
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   Peptide
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REGISTRATION NUMBER: 3033
REFERENCE/DOCKET NUMBER:
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  STREET:
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  Virginia
  Application US/08455625
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  Conservative
  Shirai, Mutsunori
  Berzofsky, Jay A. Ahlers, Jeffrey D. Pendleton, C. D.
   Nara, Peter
  Birch, Stewart, Kolasch & Birch
O. Box 747
  703-205-8000
   internal
  peptide
   LYMPHOCYTES AGAINST HIV
  COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
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  US/08/455,625
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   Length 15;
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   Gaps
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  APPLICANT:
   TOPOLOGY:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
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  APPLICATION NUMBER: 08/060
FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148
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  NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-42,306
   STREET: 225
CITY: Boston
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7; Conserv
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6214347
  amino acid
   225 Franklin Street
  617/542-8906
  Berzofsky,
Ahlers, Je
Shirai, Mutsunori
VENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
VENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
VENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
   Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. David
  Shirai, Muteunori
Shirai, Multilbetrminant Peptides That Elicit
VENTION: MELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
VENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
   Nara, Peter
   Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. David
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  Fish & Richardson P.C.
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   87.2%; Score 34; DB 3; 100.0%; Pred. No. 0.84;
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  07/751,998
  US/08/455,685
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FILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/151,998
APPLICATION NUMBER: 07/751,998
FILING DATE: 28-AUG-1991
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FILING DATE: 28-AUG-1991
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FILING DATE: 28-AUG-1991
APPLICATION NUMBER: 07/148,692
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Best Local Similarity 87.5%;
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MEDIUM TYPE: Diskett
COUNTRY: US
ZIP: 02110-2804
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MEDIUM TYPE: Diskett
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TELEPHONE: 617/542-5070
  FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
  CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C
STREET: 225 Franklin Street
  APPLICANT:
  NUMBER OF SEQUENCES:
   APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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  1 RAFVTICK 8
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   RAFYTIGK 15
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  Application US/08455685
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   617/542-8906
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   Pendleton, C. David
  Berzofsky, Jay A.
Ahlers, Jeffrey D.
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Diskette
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FILING DATE: 26-JAN-1988
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NAME: Beattle, Ingrid A.
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SOFTWARE: FastSEQ for Windows Version 2.0
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FILING DATE: 06-MAR-1992
FILING TATE: 07/751,998
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TOPOLOGY: linear
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   APPLICATION NUMBER: FILING DATE: 31-MA
   Local Similarity conserv
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  COMPUTER:
  STATE:
  STREET: 225 F
               APPLICATION NUMBER: 08/0
FILING DATE: 14-MAY-1993
  LENGTH:
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  ADDRESSEE:
   INFORMATION:
  1 RAFVTIGK 8
   RAFVAIGK 15
  200154
  M
  Application US/08455685
   225 Franklin Street
  15 amino acide
  Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. David
  Conservative
   Shirai, Mutsunori
  Nara, Peter
   IBM Compatible
  Fish & Richardson P.C.
   14-MAY-1993
  31-MAY-1995
  87.2%;
  MULTIDETERMINANT PEPTIDES THAT ELICIT HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
   07/148,692
                              08/060,988
  08/060,988
07/847,311
  US/08/455,685
  08830/022003
  Score 34; DB
Pred. No. 0.84
0; Mismatches
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   Length 15;
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  Gaps
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밁
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  US-08-060-988A-16
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Best Local Similarity 87.5%;
Matches 7; Conservative
   Sequence 16, Application US/08060988A Patent No. 6294322
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-WAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-WAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/151,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-198
ATTORNEY/AGENT INFORMATION:
NAME: Beattle, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 0830/022001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
  TELEX: 200154
INFORMATION FOR SEQ ID NO:
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  SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
  NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/022003
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
   APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 48
   FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
  STREET: 2. STREET: 2. STREET: MA
  CORRESPONDENCE ADDRESS
   TELLEFAX: 0. TELEFAX: 200154
   COMPUTER: IBM CONTROL OPERATING SYSTEM: SOFTWARE: FastSE(
   COUNTRY: US
ZIP: 02110-2804
   ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
  APPLICATION NUMBER:
   FILING DATE: 06-MAI APPLICATION NUMBER:
   8 RAFVITCK 15
   1 RAFVTICK 8
   617/542-8906
  Berzofsky, Jay A.
Ahlers, Jeffrey I
   Nara, Peter
   Pendleton, C. David
  IBM Compatible
SYSTEM: Windows95
   06-MAR-1992
  29-AUG-1991
  07/148,692
   07/751,998
   Score 34; DB 3; Length 15; Pred. No. 0.84; 0; Mismatches 1; Indels
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US-08-060-988A-19
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LENGTH: 15 aming
Type
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OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/060,988A
FILING DATE: 14-MAY-1993
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APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/151,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
ETILNG DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
   Query Match
  FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 08830/
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
   TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
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ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
   APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES
TITLE OF INVENTION: THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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  MOLECULE TYPE: peptide
  APPLICANT:
  APPLICANT:
   Local Similarity nes 7; Conserv
  TOPOLOGY:
  CITY: Boston
  COUNTRY:
  2 AFVTIGK 8
   9, Application US/08060988A
6294322
  amino acid
  ¥
  เร
  Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. David
  Nara, Peter
   Conservative
   87.2%;
  87.2%; Score 34;
   19:
   16:
  08830/022001
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   Score 34; DB 3; Pred. No. 0.84;
   Mismatches
  DB 3;
  Length 15;
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   Indels
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   Gaps
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Matches

Best Local Similarity 87.5%; Matches 7; Conservative

0

Pred. No. 0.84; 0; Mismatches

1:

Indels

0

Gaps

0

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; Sequence 21, Application US/08060988A
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US-08-060-988A-21
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INFORMATION FOR SEQ ID NO:
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TITLE OF INVENTION:
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LENGTH: 15 amino acids
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PRIOR APPLICATION DATA: 07/847,311
APPLICATION NUMBER: 07/847,311
   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
  APPLICATION NUMBER: 07/84:
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/75:
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/14:
  SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
  TELEPHONE: 617/542-5070
   APPLICATION NUMBER: 07/148,692 FILING DATE: 26-JAN-1988 ATTORNEY/AGENT INFORMATION:
  CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
   TELEFAX: bi.,
TELEFAX: bi.,
200154
  TUMBER OF SEQUENCES:
   APPLICANT:
  NAME: Beattie, Ingrid A. REGISTRATION NUMBER: P-4 REFERENCE/DOCKET NUMBER:
  COMPUTER: IBM CONTROL OPERATING SYSTEM:
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   TELEPHONE: 61//542-8906
   STREET:
   COUNTRY: US
ZIP: 02110-2804
   Similarity 7; Conserv
   Boston
  RAFYTIGK 15
  RAFVAIGK 15
   ¥
   RAFVTIGK 8
  Application US/08060988A
  225 Franklin Street
   Conservative
  Nara, Peter
   Shirai, Mutsunori
  Berzofsky, Jay A.
Ahlers, Jeffrey D.
   Pendleton, C. David
  IBM Compatible
  MULTIDETERMINANT PEPTIDES
THAT ELLCIT
HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
  87.2%;
87.5%;
  Windows95
  07/751,998
  P-42,306
  08830/022001
   0,
   Score 34; DB 3;
Pred. No. 0.84;
0; Mismatches
  Length 15;
   Indels
   <u>.</u>
   Gaps
   0
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Query Match
Best Local Similarity
Marches 7; Conserve
   밁
   S
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   RESULT 183
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  FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
  TELEX: 200154
INFORMATION FOR SEQ ID NO:
   GENERAL INFORMATION:
  APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
   SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
   REFERENCE/DOCKET NUMBER: 08
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TELEPHONE: 617/542-5070
   OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
  APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETH
TITLE OF INVENTION: THAT ELIC
TITLE OF INVENTION: HELPER T.
TITLE OF INVENTION: NEUTRALIZ
                                 NUMBER OF SEQUENCES: 1
   APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPBUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VII
  MOLECULE TYPE:
   FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/8
FILING DATE: 06-MAR-1992
  COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
  CORRESPONDENCE ADDRESS
   NUMBER OF SEQUENCES:
  APPLICATION NUMBER:
   COMPUTER: IBM CON
OPERATING SYSTEM:
   COUNTRY: US
ZIP: 02110-2804
  ADDRESSEE: KIBH & ... STREET: 225 Franklin Street
 STREET:
                    ADDRESSEE:
   TOPOLOGY:
  TELEFAX:
  amino acid
   RAFVTIGK 8
   RAFVITGK 15
   MA

    Robbins, Berliner & Carson
    Rorth Figueroa Street, Suite 500

  617/542-8906
   15 amino acids
  Conservative
   Nara, Peter
   Ahlers, Jeffrey D.
Pendleton, C. David
  Berzofsky,
  Douvas, Angeline
   IBM Compatible
   Fish & Richardson P.C.
  peptide
  MULTIDETERMINANT PEPTIDES
THAT ELICIT
HELDER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
   87.2%;
87.5%;
  121
   07/847,311
  US/08/060,988A
  21:
  0,
  08830/022001
   Score 34; DB 3;
Pred. No. 0.84;
  Mismatches
  CLUSTER VIRUS INFECTIONS
  Length 15;
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Los Angeles

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APPLICATION NUMBER: PCT/US94/05142
FILING DATE: 13-MAY-1994
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
  PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/029
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:

NAME: Spitals, John P.

REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
   TELEFAX: (213) 977-1003 INFORMATION FOR SEQ ID NO:
   TITLE OF INVENTION: OF TITLE OF INVENTION: ITTILE OF INVENTION: ITTILE OF SEQUENCES:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   SEQUENCE CHARACTERISTICS:
   SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
   MOLECULE TYPE: peptide
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
   STATE: Virginia
   COUNTRY: USA
  TYPE: amino acid TOPOLOGY: linear
  COUNTRY:
  STATE: California
  ADDRESSEE:
  LENGTH:
   9
  1 RAFVTIG 7
  22040-0747
   90012
   15 amino acids
  U.S.A.
   Conservative
  Birch, Stewart, Kolasch & Birch
O. Box 747
   COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC TLYMPHOCYTES AGAINST HIV
   87.2%; Score 34; DB 5; Length 15; 100.0%; Pred. No. 0.84; tive 0; Mismatches 0; Indels
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   1920-331
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LOCATION: 1..15
; OTHER INFORMATION: /;
; OTHER INFORMATION: /;
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Best Local Similarity
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   ঠ
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TELEPHONE: 703-205-8050
TELEPAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 19:
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APPLICATION NUMBER: US 08/0
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORWATION:
NAME: SVERBSON, LECHART R.
REGISTRATION NUMBER: 30330
  GENERAL INFORMATION:
  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
   TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: pepti
FRAGMENT TYPE: inter
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LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
   MOLECULE TYPE:
FRAGMENT TYPE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
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  FEATURE:
   ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
CITY: Falls Church
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OTHER INFORMATION: /note= "p18-8, see Table V"
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   Virginia
  Application PC/TUS9405142
  703-205-8050
  Conservative
   peptide
internal
  13-MAY-1994
   internal
   peptide
  COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC TLYMPHOCYTES AGAINST HIV
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100.0%; Pred. No.
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   PCT/US94/05142
   30330
  16:
   1173-434P
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  Mismatches
  DB 5;
  Length 15
  <u>,</u>
  Gaps
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Query Match Best Local Similarity

87.2%;

Length 15

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  RESULT 186
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  RESULT 187
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              Sequence 21, Application PC/TUS9405142 GENERAL INFORMATION:
APPLICANT:
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Matches
  Matches
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SENERAL INFORMATION:
   INFORMATION FOR SEQ ID NO:
   ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Re-lease #1.0, Version #1.25
CURRENT APPLICATION DATA:
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LENGTH: 15 amino acids
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  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,988
FILING DATE: 14-MAY-1993
   TITLE OF INVENTION:
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TITLE OF INVENTION:
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  LOCATION: 1..15
OTHER INFORMATION:
OTHER INFORMATION:
   TELEPHONE: /UJ -8050
  TYPE: amino acid
TOPOLOGY: linear
  NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330
   STREET: P.O. Box 7.
CITY: Falls Church
  STREET:
   NAME/KEY:
   REFERENCE/DOCKET NUMBER:
  COUNTRY:
  ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. Box 747
   œ
   1 RAFVTIGK 8
  1 RAFVTIGK 8
   Similarity 7; Conserv
   RAFVAIGK 15
   Virginia
  Application PC/TUS9405142
  Conservative
   Conservative
  Peptide
   peptide
internal
   13-MAY-1994
  87.2%;
87.5%;
   LYMPHOCYTES AGAINST HIV
COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
   COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
  /label= peptide
/note= "p18-12,
  PCT/US94/05142
  30330
ED: 1173-434P
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Pred. No. 0.84
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  Mismatches
   see Table V"
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   Indels
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   0;
  Gaps
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   0
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COMPUTER READABLE FORM:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   NUMBER OF SEQUENCES: 1
  APPLICANT: SIA, Charles D.Y
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem
  SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
   NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330 REFERENCE/DOCKET NUMBER: 11 TELECOMMUNICATION INFORMATION:
  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
  TITLE OF INVENTION: ELTITLE OF INVENTION: LY.
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
   MOLECULE TYPE:
FRAGMENT TYPE:
  STREET: Suite
CITY: Toronto
STATE: Ontario
  Local Similarity tes 7; Conserv
  APPLICATION NUMBER: FILING DATE: 13-MAN CLASSIFICATION:
  LOCATION: 1..15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-13, see Table V"
   TELEPHONE: 703-205-8050
  STREET: P.O. Box 7
CITY: Falls Church
  COUNTRY:
  COUNTRY:
   ADDRESSEE:
   NAME/KEY: Peptide
  TYPE: amino acid TOPOLOGY: linear
  ADDRESSEE:
   ENGTH:
   1 RAFVTICK 8
   22040-0747
  Virginia
  Ontario
   Application US/08257528B
   Suite 701,
  Birch, Ste
   Canada
   Conservative
  Sim & McBurney
   703-205-8000
  peptide
internal
   13-MAY-1994
   Charles D.Y.
   87.2%;
87.5%;
  ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T LYMPHOCYTES AGAINST HIV
   Tandem Synthetic HIV-1 Peptides
   Stewart, Kolasch & Birch
  101
  330 University Avenue
  30330
1173-434P
   US 08/060,988
   PCT/US94/05142
   Score 34; DB
Pred. No. 0.84
0; Mismatches
  DB 5;
  Length 15
   Indels
   <u>,,</u>
   Gaps
   0,
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片 S

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무
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   Sequence 35, Application US/08460602A Patent No. 5759769
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  TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1163
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 35:
TELEPHONE: (416) 595-1155
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 35:
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APPLICATION NUMBER: 06
FILING DATE: 09-JUN-15
CLASSIFICATION: 424
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FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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LENGTH: 17 amino acids
TYPE: amino acid
   ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10.
TELECOMMUNICATION INFORMATION:
  SOFTWARE: PatentIn Release #1.0, Version #1.25
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   87.2%; Score 34; DB 1; Local Similarity 100.0%; Pred. No. 0.95; 1es 7; Conservative 0; Mismatcher
  APPLICATION NUMBER: US/01
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
  CITY: Toronto
STATE: Ontario
  STREET:
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REFERENCE/DOCKET NUMBER: 1038-336 MIS:jb
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  ADDRESSEE:
   11 RAFVTIG 17
   1 RAFVTIG 7
  INVENTION:
  STEWART, MICHAEL I.
  Suite 701, 330 University Avenue
  Canada
  SIA, Charles D.Y.
CHONG, Pele
KLEIN, Michel H.
   linear
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   09-JUN-1993
   single
  Tandem Synthetic HIV-1 Peptides
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  US/08/257,528B
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   Gaps
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밁
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   US-08-463-966A-35
  US-08-460-602A-35
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   RESULT 190
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   Sequence 35, Application US/08463966A Patent No. 5795955
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  APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REGISTRATION NUMBER: 1038-448
  TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 35
   GENERAL INFORMATION:
   SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
  REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
   CORRESPONDENCE ADDRESS:
  NUMBER OF SEQUENCES:
   APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H
  TITLE OF INVENTION:
  APPLICANT:
  y Match 87.2%;
Local Similarity 100.0%;
hes 7; Conservative
  APPLICATION NUMBER: US/08/463,966A FILING DATE: 05-JUN-1995 CLASSIFICATION: 424
   STRANDEDNESS:
TOPOLOGY: lir
   TYPE: amino acid
STRANDEDNESS: sin
   TOPOLOGY:
  CITY: Toronto
  COUNTRY:
  STREET:
11 RAFVTIG 17
   11 RAFVTIG 17
  1 Similarity 100
    7; Conservative
  rRY: Canada
M5G 1R7
                               1 RAFVTIG 7
  1 RAFVTIG 7
   Ontario
  E: Sim & McBurney
Suite 701, 330 University Avenue
   linear
   linear
  single
  Charles D.Y.
   single
  87.2%; Score 34; DB 1; 100.0%; Pred. No. 0.95; tive 0; Mismatches
  Tandem Synthetic HIV-1 Peptides
  101
  Score 34; DB 1; Length 17; Pred. No. 0.95; 0; Mismatches 0; Indels
   1038-487 MIS:jb
   Length 17;
  Indels
  <u>,</u>
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  Gaps
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US-08-465-217A-35
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Patent No. 5817754
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,217A
FILING DATE: 05-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
DATA: 08/257,528
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  SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
  CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
   REFERENCE/DOCKET NUMBER: 1038-486 MIS:jb TELECOMMUNICATION INFORMATION:
   ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,9
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
   CORRESPONDENCE ADDRESS
   TITLE OF INVENTION:
   APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
   STREET: STREET: Toronto
  CORRESPONDENCE ADDRESS
  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
   APPLICATION NUMBER: 08/2 FILING DATE: 09-JUN-1994
   COUNTRY: Canada
ZIP: M5G 1R7
  STRANDEDNESS:
  TELEPHONE:
                                 STREET:
  LE OF INVENTION:
  DDRESSEE:
   DDRESSEE:
   35, Application US/08465217A
5. 5800822
   11 RAFVTIG 17
   1 RAFVTIG 7
                 Toronto
   Ontario
Ontario
   E: (416)
(416) 59
   E: Sim & McBurney
Suite 701, 330 University Avenue
                              B: Sim & McBurney
Suite 701, 330 University Avenue
  SIA, Charles D.Y
CHONG, Pele
KLEIN, Michel H.
  SIA, Charles D.Y.
  PatentIn.Release #1.0, Version #1.25
   linear
   Floppy disk
   Charles D.Y.
   single
   595-1163
   Tandem Synthetic HIV-1 Peptides
  87.2%; Score 34; DB 1; Length 17; 100.0%; Pred. No. 0.95;
  595-1155
  Tandem Synthetic HIV-1 Peptides
   101
   24,973
  Mismatches
  0; Indels
  0
  Gaps
  0
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US-08-462-507A-35
   US-08-464-329A-35
  Best Loc
Matches
  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IEM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/462,507A

FILING DATE: 05-UN-1995

CLASSIFICATION: 424

PRIOR APPLICATION DATA:

CLASSIFICATION DATA:

OS COMPANY CON CONTROL OF CONTROL O
   Patent No.
   Query Match
   TELEFAX: (416) 595-1163 INFORMATION FOR SEQ ID NO:
  GENERAL INFORMATION:
  CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-449 MIS:jb
TELECOMMUNICATION INFORMATION:
  SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
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APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
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MEDIUM TYPE: Floppy
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APPLICATION NUMBER: US,
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
   APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
   COUNTRY! Carrio
  STREET: Suite
CITY: Toronto
STATE: Ontario
  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Palana
  Local Similarity
es 7; Conserv
  FILING DATE:
  CLASSIFICATION:
   TOPOLOGY:
   STRANDEDNESS:
   APPLICATION NUMBER:
   ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
  ELEPHONE:
   FILING DATE: 09-JUN-1993
  APPLICATION NUMBER:
   11 RAFVTIG 17
  M5G 1R7
  1 RAFVTIG 7
  Application US/08462507A
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  (416)
   PatentIn Release #1.0, Version #1.25
  Canada
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  (416)
   05-JUN-1995
  Charles D.Y
   single
   595-1155
  Tandem Synthetic HIV-1 Peptides
   100.0%;
   101
   08/073,378
   08/257,528
  US/08/464,329A
   35:
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   Gaps
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FILING DATE: 06-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
PRIOR DATE: 09-JUN-1993
CLASSIFICATION: 424
PRIOR DATE: 09-JUN-1993
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   Sequence 35, Application US/08467881A Patent No. 5951986
  COUNTRY: CANADA
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, V
  GENERAL INFORMATION:
  TELEFAX: (416) 595-1163 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                      TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
  tent No.
   TYPE: amino acids
STRANDEDNESS: single
TOPOLOGY: li-
   REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-488 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
  ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I
REGISTRATION NUMBER: 24,
   NUMBER OF SEQUENCES: 1
  APPLICANT:
  REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-451 MIS:jb
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
  ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I
  STREET: our
CTTY: Toronto
   CLASSIFICATION:
   APPLICATION NUMBER: 08/073,378
   ADDRESSEE:
   11 RAFVTIG 17
  1 RAFVTIG 7
: 17 amino acids amino acid
   INVENTION:
  Ontario
  SIA, Charles U.x
CHONG, Pele
KLEIN, Michel H.
   E: Sim & McBurney
Suite 701, 330 University Avenue
   09-JUN-1993
  Charles D.Y.
   Tandem Synthetic HIV-1 Peptides
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RESULT 195
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  ; TOPOLOGY: linear US-08-467-881A-35
  문
   S
  PCT-US92-06688-13
   Query Match
Best Local Similarity lov.
Thickes 7; Conservative
  Sequence
  Matches
   Sequence 13, Applicat GENERAL INFORMATION:
   Patent No.
  Best Local Similarity Matches 7; Conserv
   Query Match
   -08-455-625-18
                                GENERAL INFORMATION:
APPLICANT: Berzof
  TELEX: 200154
INFORMATION FOR SEQ ID NO:
   CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 744,26
PILING DATE: 13 AUGUST 199
ATTORNEY/AGENT INFORMATION:
NAME: Paul T. Clark
  SEQUENCE CHARACTERISTICS:
LENGTH: 17
   REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00231/052W01
TELECOMMUNICATION INFORMATION:
   COMPUTER: IBM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: IBM P.C. DOS (Vers)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
  NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
   APPLICANT: REPLIGEN CORPORATION
APPLICANT: THE ROCKEFELLER UNIVERSITY
TITLE OF INVENTION: MULTIPLE ANTIGEN PEPTIDES FOR USE AS
TITLE OF INVENTION: VACCINES
  TELEFAX: 10-
TELEFAX: 200154
  CITY: Boston
STATE: Massac
   TYPE: AMINO ACID
  TOPOLOGY:
  TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
   APPLICATION NUMBER: FILING DATE: 19920
   ZIP: 02110-2804
  COUNTRY:
   ADDRESSEE:
   11 RAFVTIG 17
  1 RAFVTIG 7
  1 RAFVTIG 7
   Massachusetts
  Application US/08455625
   Application PC/TUS9206688
  225 Franklin Street
Berzofsky, Jay A. Ahlers, Jeffrey D. Pendleton, C. D.
  Conservative
  linear
   Fish & Richardson
   19920811
  87.2%; 5--
/ 100.0%; Pred. No.
/ 0; Mismatches
  100.0%;
  87.2%; Score 34; DB 5; 100.0%; Pred. No. 0.95;
   744,281
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  DB 2;
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US-08-455-625-22
  밁
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   GENERAL INFORMATION:
   PILING DATE:
CLASSIFICATION: 435
CLASSIFICATION DATA;
PRIOR APPLICATION NUMBER: US 08/00
APPLICATION NUMBER: US 08/00
APPLICATION NUMBER: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 30330
   TELEFAX: 703-205-8050
   APPLICANT: Shirai, I
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
   APPLICANT:
  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
  APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT
TITLE OF INVENTION: ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T
TITLE OF INVENTION: LYMPHOCYTES AGAINST HIV
  APPLICANT:
   FEATURE:
NAME/KEY: Peptide
  SEQUENCE CHARACTERISTICS:
  NUMBER OF SEQUENCES: 3
  CORRESPONDENCE ADDRESS:
   TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
PRAGMENT TYPE: internal
  TELECOMMUNICATION INFORMATION: TELEPHONE: 703-205-8000
   TUMBER OF SEQUENCES:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: P.O. BOX 747
CITY: Falls Church
STATE: Virginia
  REGISTRATION NUMBER: 30330
REFERENCE/DOCKET NUMBER: 1173-434P
  APPLICATION NUMBER:
  STREET: P.O. Box 7.
  ADDRESSEE: Birch, St
STREET: P.O. Box 747
  COUNTRY: USA
   TELEPHONE:
   ENGTH:
  ထ
   22040-0747
   RAIVTIGK 15
  RAFYTIGK 8
   Virginia
  15 amino acids
  Berzofsky, Jay A.
Ahlers, Jeffrey D.
   Conservative
  Nara, Peter
  Pendleton, C. D.
   Mutsunori
   COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC TLYMPHOCYTES AGAINST HIV
  84.6%;
87.5%;
   Stewart, Kolasch & Birch
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/note= "p18-10, see Table V"
   US 08/060,988
  US/08/455,625
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Pred. No. 1.4;
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   DB 2;
   1; Indels
   Length 15;
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   Gaps
   0;
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멹
  Query Match
Best Local Similarity
Thes 7; Conserve
  Ş
   US-08-455-685-18
   RESULT 198
  US-08-455-625-22
  Sequence 18, App. No. 621434
  CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 08/060,988
APPLICATION NUMBER: 193
ATTORNEY/AGENT INFORMATION:
NAME: Svensson, Leonard R.
REGISTRATION NUMBER: 3030
REGISTRATION NUMBER: 3173-434P
   GENERAL INFORMATION:
   TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO:
 COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: Fast-SEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
  COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
  COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
   APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDET
TITLE OF INVENTION: HELPER T-
TITLE OF INVENTION: NEUTRALIZ
   APPLICANT:
  TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
   SEQUENCE CHARACTERISTICS LENGTH: 15 amino acid
   REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
   APPLICANT:
  FEATURE:
  NUMBER OF SEQUENCES:
   APPLICANT:
   APPLICANT:
  NAME/KEY: Peptide LOCATION: 1..15 OTHER INFORMATION: OTHER INFORMATION:
   TELEPHONE: 703-205-8050
   FILING DATE:
   APPLICATION NUMBER:
   CITY: Boston
FILING DATE:
   æ
  1 RAFVTICK 8
  ¥
   RAFVTIAK 15
   Application US/08455685
  15 amino acids
   Nara, Peter
   Berzofsky, Jay -- Day -- Day
  Conservative
   Pendleton, C. David
   703-205-8000
31-MAY-1995
   MULTIDETERMINANT PEPTIDES THAT ELICIT HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
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  /label= peptide
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   Gaps
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; LENGTH: 15 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-455-685-18
   US-08-455-685-22
; Sequence 22, A
  Query Match 84.6%;
Best Local Similarity 87.5%;
Matches 7; Conservative
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOPTWARE: PastSQC for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,685
PILING DATE: 31-MAY-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/060,988
PILING DATE: 14-MAY-1993
APPLICATION NUMBER: 07/847,311
PILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/51,998
PILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
PILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
PILING DATE: 26-JAN-1988
  stent No. 6214347
GENERAL INFORMATION:
  TELEX: 200154
INFORMATION FOR SEQ ID NO:
   FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
   ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
  APPLICANT: Nara, Peter
APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETERMINANT PEPTIDES THAT ELICIT
TITLE OF INVENTION: HELPER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
NUMBER OF SEQUENCES: 40
  TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
  PRIOR APPLICATION DATA:
APPLICATION NUMBER: . 08/0
FILING DATE: 14-MAY-1993
  ADUAL STREET: LE BOSTON MA
   NAME: Beattie, Ingrid A
REGISTRATION NUMBER: P-4
REFERENCE/DOCKET NUMBER:
   COUNTRY:
   APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILING DATE: 29-AUG-1991
   1 RAFVTICK 8
  RAIVTICK 15
  Application US/08455685
  225 Franklin Street
   S
   Pendleton, C. David
   Berzofsky, Jay A.
Ahlers, Jeffrey D.
  08/060,988
  P-42,306
  Score 33; DB 3; Length 15;
Pred. No. 1.4;
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   08830/022003
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US-08-060-988A-18
  US-08-455-685-22
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Best Local Similarity 87.5%;
Matches 7; Conservative
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PTILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION UNMER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/751,998
FILLING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILLING DATE: 26-JAN-1988
ATTORNEY/AGENT INFORMATION:
NAME: Beattie, Ingrid A.
REGISTRATION NUMBER: P-42,306
REGERENCE/DOCKET NUMBER: 08830/0:
TELECOMMUNICATION INFORMATION:
   TRLEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
  TELEX: 200154
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: PastSEQ for Windows Version
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LENGTH: 15 amino acids
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
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TITLE OF INVENTION: HILDER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: HELDER T-LYMPHOCYTE, CYTOTOXIC T LYMPHOCYTE AND
TITLE OF INVENTION: NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
  NUMBER OF SEQUENCES: 4
  APPLICANT: Berzofsky, Jay A. APPLICANT: Ahlers, Jeffrey D.
  CITY: Boston
STATE: MA
  ADDRESSEB: Fisn a ADDRESSEB: Fisn a ADDRESSEB: STREET: 225 Franklin Street
   TELEFAX:
  TELEPHONE:
   TOPOLOGY:
   COUNTRY:
   TELEFAX:
  REGISTRATION NUMBER:
  629432
   amino acid

GY: linear
  RAFVTIGK 8
  Beattie, Ingrid A
   617/542-8906
   617/542-8906
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  Nara, Peter
Shirai, Mutsunori
  Pendleton, C. David
  Fish & Richardson P.C.
   MBER: US/08/060,988A
14-MAY-1993
  P-42,306
  18:
   08830/022001
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  Score 33; DB 3;
Pred. No. 1.4;
  Length 15
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   Gaps
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밁
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Best Local Similarity
7; Conserve
                           Ś
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   US-08-060-988A-22
  RESULT 201
US-08-060-988A-22
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  Query Match
Best Local Similarity 87.3
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OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U5/08/060,988A
FILING DATE: 14-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/847,311
FILING DATE: 06-MAR-1992
APPLICATION NUMBER: 07/51,998
FILING DATE: 29-AUG-1991
APPLICATION NUMBER: 07/148,692
FILING DATE: 26-DAN-1989
ATTORNEY/AGENT INFORMATION:
NAME: BESTIF OF TOTAL
   Patent No.
   INFORMATION FOR SEQ ID NO:
   GENERAL INFORMATION:
   SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
  TOPOLOGY: li
  TELEPHONE: 617/542-5070
  COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
  APPLICANT: Shirai, Mutsunori
TITLE OF INVENTION: MULTIDETI
TITLE OF INVENTION: THAT ELIC
TITLE OF INVENTION: HELPER T.
   APPLICANT:
   APPLICANT:
  APPLICANT:
  NUMBER OF SEQUENCES:
  TELEFAX:
   TELEPHONE:
   REGISTRATION NUMBER: P-42,306
REFERENCE/DOCKET NUMBER: 0883
   COUNTRY:
  02110-2804
   Boston
  Beattie, Ingrid
                            RAFVTIGK 8
RAFVTIAK 15
   RAIVTICK 15
  RAFVTICK 8
  3
   Application US/08060988A
  617/542-8906
  SD
   Conservative
   Nara, Peter
  Berzofsky, Jay A.
Ahlers, Jeffrey D.
Pendleton, C. David
   linear
   peptide
  84.6%;
  84.6%;
   MULTIDETERMINANT PEPTIDES THAT ELICIT HELPER T-LYMPHOCYTE, CYTO
  NEUTRALIZING ANTIBODY RESPONSES AGAINST HIV-1
   0
   08830/022001
   0
  Score 33; DB 3;
Pred. No. 1.4;
  Score 33; DB
Pred. No. 1.4;
   Mismatches
  DB 3;
   CYTOTOXIC T LYMPHOCYTE AND
   Length 15;
  Length 15;
   Indels
  0
   °,
  Gaps
   Gaps
  0
   0
```

```
Ś
   RESULT 203
PCT-US94-05142-22
  밁
  PCT-US94-05142-18
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  Sequence 22, Application PC/TUS9405142 GENERAL INFORMATION:
   Matches
   Query Match
  ZIP: 22040-0747

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTMARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US94/05142

FILING DATE: 13-MAY-1994

CLASSIFICATION:
   Sequence 18, Applicat GENERAL INFORMATION:
   TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
   APPLICANT:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
   APPLICANT:
APPLICANT:
CITILE OF INVENTION: CITILE OF INVENTION: I
   TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: pepti
FRAGMENT TYPE: inter
  NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
   TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US C
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
   NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
  Local
   OTHER INFORMATION: OTHER INFORMATION:
                ADDRESSEE: Birch, Stewart, Kolasch & Birch
   LOCATION:
   NAME/KEY:
  REFERENCE/DOCKET NUMBER:
   NAME: Svensson, Leonard R. REGISTRATION NUMBER: 3033(
   ADDRESSEE: Birch, St. STREET: P.O. Box 747 CITY: Falls Church
   COUNTRY: USA
  ENGTH:
  1 RAFVTIGK 8
   Similarity 7; Conserv
: P.O. Box 7
  Virginia
  Application PC/TUS9405142
  15 amino acids
   Conservative
   Peptide
   internal
   peptide
   COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC TLYMPHOCYTES AGAINST HIV
  84.6%;
  Stewart, Kolasch & Birch
  COMPOSITE SYNTHETIC PEPTIDE CONSTRUCT ELICITING NEUTRALIZING ANTIBODIES AND CYTOTOXIC T LYMPHOCYTES AGAINST HIV
   /label= peptide
/note= "p18-10,
  US 08/060,988
   30330
  1173-434P
  Score 33; DB
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   Mismatches
   see Table
   DB 5;
   1; Indels
   Length 15;
   0
   0
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Virginia

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ફ
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   US-08-257-528B-51
  RESULT 204
   PCT-US94-05142-22
   Query Match 84.6%;
Best Local Similarity 87.5%;
Matches 7; Conservative
  Sequence 51, Appli
Patent No. 5639854
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  INFORMATION FOR SEQ ID NO:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/257,528B
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
   APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem S
NUMBER OF SEQUENCES: 101
  SEQUENCE CHARACTERISTICS
   SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compactible
OPERATING SYSTEM: PC-DOS/MS-DOS
  CORRESPONDENCE ADDRESS
   MOLECULE TYPE:
FRAGMENT TYPE:
  TELECOMMUNICATION INFORMATION: TELEPHONE: 703-205-8000
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 14-MAY-1993
ATTORNEY/AGENT INFORMATION:
  STATE: V
   NAME/KEY: Peptide
LOCATION: 1.15
OTHER INFORMATION: /label= peptide
OTHER INFORMATION: /note= "p18-14, see Table V"
  ZIP:
   STREET: Suite
CITY: Toronto
   APPLICATION NUMBER: FILING DATE: 13-MA' CLASSIFICATION:
   COUNTRY:
   STATE:
   NAME: Svensson, Leonard R. REGISTRATION NUMBER: 30330 REFERENCE/DOCKET NUMBER: 1
   TELEFAX:
  TELEPHONE:
  POPOLOGY:
   ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
   ENGTH:
  M5G 1R7
   22040-0747
  amino acid
XGY: linear
   RAFVTICK 8
  RAFVTIAK 15
   Ontario
   Application US/08257528B
   15 amino acids
   703-205-8050
   Canada
   peptide
internal
  13-MAY-1994
   Tandem Synthetic HIV-1 Peptides
  US 08/060,988
  PCT/US94/05142
  30330
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Pred. No. 1.4;
   1173-434P
  DB 5; Length 15
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  US-08-257-528B-51
  COUNTRY: Canada

ZIP: MSG 1R7

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/460,602A

FILING DATE: 02-JUN-1995

CURSETTETATION. 476
Query Match
Best Local Similarity
  Sequence 51, Application US/08460602A Patent No. 5759769
   Matches
   Query Match
Best Local Similarity
  TELEPHONE: (416) 595-111
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO:
   TELEPHONE: (416) 595-11:
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
  GENERAL INFORMATION:
   ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
   SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
   REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/0:
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
  NUMBER OF SEQUENCES: 1
  CLASSIFICATION: 424
PRIOR APPLICATION DATA:
   TITLE OF INVENTION:
   APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
  APPLICANT:
   TELECOMMUNICATION INFORMATION: TELEPHONE: (416) 595-1155
  APPLICATION NUMBER: FILING DATE: 09-JUN CLASSIFICATION: 424
  STREET: Suite CITY: Toronto
STATE: Ontario
  TOPOLOGY: 11
  ADDRESSEE:
   TOPOLOGY:
  REFERENCE/DOCKET NUMBER: 1038-336 MIS:jb
   REGISTRATION NUMBER:
   ENGTH:
   14 RAFYTIG 20
  1 RAFVTIG 7
   6
   H: 20 amino acids amino acid
   Suite 701, 330 University Avenue
  74.4%;
nilarity 85.7%;
Conservative
   linear
   linear
  Sim & McBurney
   (416) 595-1155
   09-JUN-1994
  Charles D.Y.
   Tandem Synthetic HIV-1 Peptides
74.4%;
85.7%;
  08/073,378
   08/257,528
  24,973
   51:
   1038-450 MIS:jb
Score 29;
Pred. No.
   Score 29; DB Pred. No. 13;
   Mismatches
 DB 1;
13;
   1;
   Length 20
               Length 20;
   Indels
   0
   Gaps
```

Tandem Synthetic HIV-1 Peptides

```
밁
   US-08-465-217A-51
   RESULT 207
   밁
  US-08-463-966A-51
   US-08-463-966A-51
  RESULT 206
  NAME: STEWART, MICHAEL I.

REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEPAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
  Sequence 51, Application US/08465217A
Patent No. 5800822
GENERAL INFORMATION:
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   Query Match
Best Local
  Matches
   Sequence 51, Application US/08463966A Patent No. 5795955
  GENERAL INFORMATION:
                APPLICANT:
   FILING DATE: 05-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
FILING DATE: 09-JUN-1994
   APPLICANT:
  ELLING 424
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/073,378
   APPLICATION NUMBER: 08/0:
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
  SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
   NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
   APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
   APPLICANT:
  STREET: Surrice Toronto
   APPLICATION NUMBER:
  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
   TLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
   COUNTRY:
   ocal Similarity
   14 RAFYTIG 20
   1 RAFVTIG 7
  14 RAFYTIG 20
  M5G 1R7
  6
  Ontario
  RAFVTIG 7
   Suite 701, 330 University Avenue
SIA, Charles D.Y. CHONG, Pele KLEIN, Michel H.
   Conservative
   Canada
  Conservative
   Sim & McBurney
   Charles D.Y.
   74.48;
85.78;
  101
  US/08/463,966A
   0,
  1038-487 MIS:jb
   Score 29; DB
Pred. No. 13;
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   Mismatches
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  ۲.
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   0,
   Gaps
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   0;
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Query Match
Best Local Similarity
Matches 6; Conserve
  US-08-465-217A-51
  Sequence 51,
  Patent No.
  -08-464-329A-51
   GENERAL INFORMATION:
APPLICANT: SIA, Charles D.Y.
APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides
NUMBER OF SEQUENCES: 101
  CLASSIFICATION: 424
PRIOR APPLICATION DATA:
RAPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I
REGISTRATION NUMBER: 24,973
   INFORMATION FOR SEQ ID NO:
  ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,217A
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
  SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
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   CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNIMBER: 08/257,528
   TELECOMMUNICATION INFORMATION:
  TITLE OF INVENTION: TO NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
URRENT APPLICATION DATA:
  CITY: Toronto
  STREET:
  STRANDEDNESS:
  ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
   APPLICATION NUMBER: 08/2 FILING DATE: 09-JUN-1994
   REGISTRATION NUMBER: 24,973 REFERENCE/DOCKET NUMBER: 10
  FILING DATE:
   STREET: Suite
CITY: Toronto
   TELEPHONE:
   COUNTRY:
  ADDRESSEE:
   14 RAFYTIG 20
   MSG 1R7
   1 RAFVTIG 7

    Application US/08464329A
5817754

  Ontario
   Ontario
   Conservative
   Suite 701, 330 University Avenue
   (416) 595-1163
  linear
  Sim & McBurney
   (416)
  05-JUN-1995
  single
   74.4%;
85.7%;
   595-1155
   51:
   1038-486 MIS:jb
   Score 29; DB 1;
Pred. No. 13;
   Length 20
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   <u>.</u>.
   Gaps
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US-08-462-507A-51
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APPLICANT: SIA, C
  TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
   equence 51,
  tent No.
   CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
CLASSIFICATION: 424
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATEN: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/462,507A
PILING DATE: 05-JUN-1995
  APPLICANT: CHONG, Pele
APPLICANT: CHONG, Michel H.
APPLICANT: KLEIN, Michel H.
APPLICANT: TANDEM Synthetic HIV-1 Peptides
ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I.
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10:
  CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,528
PILING DATE: 09-JUN-1994
   REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/073,378
FILING DATE: 09-JUN-1993
CLASSIFICATION: 424
   STREET: STREET: Onto
   ATTORNEY/AGENT INFORMATION:
NAME: STEWART, MICHAEL I
  CORRESPONDENCE ADDRESS
  PRIOR APPLICATION DATA:
  APPLICATION NUMBER: 08/2
FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
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   1 RAFVTIG 7
   6; Conservative
   Ontario
   Application US/08462507A
  E: Sim & McBurney
Suite 701, 330 University Avenue
  (416) 595-1155
  08/257,528
  1038-449 MIS:jb
 1038-451 MIS:jb
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FILING DATE: 09-JUN-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMER: 08/073,378
FILING DATE: 09-JUN-1993
   TELEFAX: (416) 595-1163 INFORMATION FOR SEQ ID NO:
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   TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 1
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   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
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LENGTH: 20 amino acids
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  SEQUENCE CHARACTERISTICS
   TELECOMMUNICATION INFORMATION
   CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
  TITLE OF INVENTION: Tandem Synthetic HIV-1 Peptides NUMBER OF SEQUENCES: 101
  CORRESPONDENCE ADDRESS
  APPLICANT: CHONG, Pele
APPLICANT: KLEIN, Michel H.
y Match 74.4%;
Local Similarity 85.7%;
hes 6; Conservative
   STRANDEDNESS:
   REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-488 MIS:jb
  FILING DATE:
  COUNTRY: Can
ZIP: M5G 1R7
  STREET: Suite
CITY: Toronto
  STRANDEDNESS:
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  TYPE: amino acid
   TELEPHONE:
  ADDRESSEE:
   TELEPHONE:
   OPOLOGY:
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  14 RAFYTIG 20
  1 RAFVIIG 7
  5951986
   amino acids
   STEWART, MICHAEL I.
  Ontario
  Application US/08467881A
  E: Sim & McBurney
Suite 701, 330 University Avenue
   . single
linear
   20 amino acids
  Canada
   linear
   (416)
   (416) 595-1155
  06-JUN-1995
  Charles D.Y.
  single
   595-1155
   51:
   Score 29; DB
Pred. No. 13;
0; Mismatches
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Pred. No. 13;
                                     DB 2; Length 20;
   Length 20;
  Indels
       Indels
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   Gaps
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RAFVTIG 7

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   US-09-430-470-24
  US-09-820-484-8
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SEQ ID NO 24
LENGTH: 10
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Sequence 8, App---
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Sequence 8, App---
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Patent No. 6562800
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EARLIER FILING DATE: 1998-10-30
NUMBER OF SEQ ID NOS: 25
   GENERAL INFORMATION:
  CURRENT APPLICATION NUMBER: US/09/430,470 CURRENT FILING DATE: 1999-10-29
  APPLICANT: MCMIllan, Minnie
TITLE OF INVENTION: THE USE OF IMMUNEOTENTIATING
TITLE OF INVENTION: FOR INDUCING IMMUNE RESPONSE
FILE REFERENCE: 13761-725
  APPLICANT: Horner, Anthony A.
TITLE OF INVENTION: Method for Increasing a Cytotoxic T
TITLE OF INVENTION: Lymphocyte Response in vivo.
FILE REFERENCE: 06510-188US1
CURRENT APPLICATION NUMBER: US/09/820,484
CURRENT FILING DATE: 2001-03-28
   PRIOR APPLICATION NUMBER: US 60/192,537
PRIOR FILING DATE: 2000-03-28
PRIOR APPLICATION NUMBER: US 60/203,567
PRIOR PILING DATE: 2000-05-11
PRIOR APPLICATION NUMBER: US 60/215,895
PRIOR FILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 8
  APPLICANT:
APPLICANT:
APPLICANT:
  SOFTWARE: FastSEQ for Windows Version 4.0 EQ ID NO 8
   TYPE: PRT
ORGANISM: Artificial Sequence
   OTHER INFORMATION: Residues 318-327 of gp120 (GenBank accession OTHER INFORMATION: number gi224364)
  ORGANISM: Human immunodeficiency virus (HIV)
  OTHER INFORMATION: HIV-1 class I-restricted gp120 peptide
  FEATURE:
   TYPE: PRT
   FEATURE:
  ENGTH: 10
  Local Similarity
   Local
   1 RAFVTI 6
   5 RAFVTI 10
  1 RAFVTI 6
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  Similarity 6; Conser
  Application US/09820484
  Cho, Hearn Jay
Richman, Douglas
   Conservative
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100.0%; Pre
   71.8%; Score 28; 100.0%; Pred. No.
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11;
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US-08-937-276A-5
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  Sequence 51, Application US/09454204A PATENT NO. 6663871
GENERAL INFORMATION:
APPLICANT: McMichael, Andrew APPLICANT: Hill, Adrian V.S.
  Matches
  Query Match
Best Local (
  Sequence 5, Application US/08937276A Patent No. 6592872
GENERAL INFORMATION:
                          APPLICANT:
  APPLICANT:
  INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
  COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARES PATENTIN Release #1.0, Vers
RAPPLICATION DATA:
APPLICATION UNMEER: US/08/937,276A
FILING DATE: 15-Sep-1997
CLASSIFICATION: CUNknown>
PRIOR APPLICATION DATA:
   APPLICATION NUMBER: US 60/025,270
FILING DATE: 17-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Kenneth A.
REGISTRATION NUMBER: 31,677
REFERENCE/DOCKET NUMBER: 015280-22
TELECOMMUNICATION INFORMATION:
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
  CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
   TITLE OF INVENTION: Targeting Antigens to the MHC Class I
Processing Pathway With an Anthrax Toxin Fusion Protein
   NUMBER OF SEQUENCES:
  APPLICANT: Klimpel, Kurt
  1 RAFVTI 6
  Similarity 6; Conserv
   STRANDEDNESS: <Unknown>
  TYPE: amino acid
  STATE: Califo
COUNTRY: USA
   RAFVTI 10
Gilbert, Sarah C.
Schneider, Jorg
Plebanski, Magdalena
   TOPOLOGY: linear
  TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
   ZIP: 94111-3834
  CITY: San Francisco
  71.8%; Score 28; ilarity 100.0%; Pred. No. Conservative 0; Mismatci
   Goletz, Theres
   Berzofsky,
  California
   Stephen H.
  Theresa J.
  Mismatches
   015280-290100US
  DB 4; Length 10;
   Version #1.30
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  <u>.</u>.
  Gaps
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PRIOR APPLICATION NUMBER: DCT/GB98/01681
PRIOR FILING DATE: 1998-06-09
PRIOR FILING DATE: 1998-06-09
PRIOR FILING DATE: 1998-06-09
NUMBER OF SEQ ID NOS: 78
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 51
LENGTH: 10
TYDE: PRT
ORGANISM: Unknown
FEATURE:
  CURRENT APPLICATION NUMBER: US/09/454,204A
CURRENT FILING DATE: 1999-12-09
PRIOR APPLICATION NUMBER: PCT/GB98/01681
PRIOR FILLING DATE: 1998-06-09
PRIOR APPLICATION NUMBER: GB 97 11957.2
PRIOR FILING DATE: 1997-06-09
NUMBER OF SEQ ID NOS: 78
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 68
  믕
밁
                                 ફ
   US-09-454-204A-68
   ) OTHER INFORMATION: CTL Epitope of HIV-1 gp120 US-09-454-204A-51
   Query Match
  Query Match
Best Local Similarity
   Matches
  Matches
   Best Local Similarity
  TITLE OF INVENTION: Methods and Reagents for Vaccination TITLE OF INVENTION: Which Generate A CD8 T Cell Immune Response FILE REFERENCE: 2907.1000-000 CURRENT APPLICATION NUMBER: US/09/454,204A CURRENT FILING DATE: 1999-12-09
  LENGTH: 10
TYPE: PRT
ORGANISM: Unknown
PEATURE:
  APPLICANT: McMichael,
APPLICANT: Hill, Adr.
APPLICANT: Gilbert,
APPLICANT: Schneider
   TITLE OF INVENTION: Methods and Reagents for Vaccination TITLE OF INVENTION: Which Generate A CD8 T Cell Immune Response FILE REFERENCE: 2907.1000-000
   equence 68,
   APPLICANT:
  APPLICANT:
  APPLICANT:
  APPLICANT:
   tent No.
  OTHER INFORMATION: CTL Peptide Epitope of HIV gag
   INFORMATION:
                                       1 RAFVTI 6
  1 RAFVTI 6
  6; Conservative
   6; Conservative
   66638.
  RAFVTI 10
 RAFVTI 10
  Blanchard, Tom
  Hanke, Tomas
Smith, Geoffrey L.
  Schneider, Jorg
Plebanski, Magdalena
   Application US/09454204A
  Gilbert, Sarah C.
  McMichael, Andrew Hill, Adrian V.S.
   Blanchard,
   Smith, Geoffrey L.
   71.8%; Score 28; DB 4; Length 10; 100.0%; Pred. No. 11;
  71.8%; Score 28; DB 4; Length 10; 100.0%; Pred. No. 11;
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   0; Indels
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RESULT 216 US-09-508-552-16

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Query Match
Best Local Similarity
Watches 6; Conserve
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; TYPE: PRT
; ORGANISM: Human immunodeficiency virus type 1
US-09-508-552-16
   US-07-847-311A-20
   SEQ.ID NO 16
  Sequence 16, Application US/09508552 Patent No. 6749856
  Patent No.
   Sequence 20, Application US/07847311A
  NUMBER OF SEQ ID NOS: SOFTWARE: Patentin V
  FILE REFERENCE: 368200PCSEQ
CURRENT APPLICATION NUMBER: US
CURRENT FILING DATE: 2000-06-
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-11
   GENERAL INFORMATION:
   PRIOR APPLICATION NUMBER: 60/0 PRIOR FILING DATE: 1998-02-17
  APPLICANT: Kelsail, Brian L.
APPLICANT: Strober, Warren
APPLICANT: THE GOVERNMENT OF THE UNITED STATES OF AMERICA,
TITLE OF INVENTION: MUCOSAL CYTOTOXIC T LYMPHOCYTE RESPONSIFILE REFERENCE: 368200PCSEQ
   APPLICANT:
  APPLICANT:
                 TELEFAX: 70
   APPLICANT:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/847,311A
FILLING DATE: 06-WAR-1992
CTRESCIPTATION.
  CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SVENBSON, Leonard R.
REGISTRATION NUMBER: 30,330
REFERENCE/DOCKET NUMBER: 1173-392P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-241-1300
  TITLE OF INVENTION: Potent Peptide for Stimulation of TITLE OF INVENTION: Cytotoxic T Lymphocytes Specific
   CORRESPONDENCE ADDRESS:
   NUMBER OF SEQUENCES:
   APPLICANT: Margulies, David H.
  APPLICANT:
   APPLICANT:
SEQUENCE CHARACTERISTICS:
   ADDRESSEE: Birch, Stewart, Kolash & Birch
STREET: 301 N. Washington
CITY: Falls Church
STATE: Virginia
  COUNTRY: USA
ZIP: 22046-0747
  INFORMATION:
  1 RAFVTI 6
  5976541
  SEQ ID NOS: 20
PatentIn Ver. 2.0
   Belyakov,
   Berzofsky,
   Berzofsky, Jay A.
Takeshita, Toshiyuki
   Conservative
                                      703-241-2848
  Shirai, Mutsunori
   Pendleton, C.D.
Koslowski, Steven
  Michael
   2000-06-
   100.0%;
  71.8%;
  60/058,523
   60/074,894
   US/09/508,552
  Score 28; DB 4; Length 10; Pred. No. 11;
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; NAME/KEY: Peptide
; LOCATION: 1..13
; OTHER INFORMATION:
; OTHER INFORMATION:
; OTHER INFORMATION:
; OTHER INFORMATION:
   RESULT 218
US-08-930-917A-14
  밁
   δ
   Query Match 71.8
Best Local Similarity 100
Matches 6; Conservative
   Sequence 14,
Patent No. 6
   APPLICATION UNMBER: PCT/CU97/00001
APPLICATION UNMBER: 17-Jan-1997
ATTORNEY/AGENT INFORMATION:
NAME: HENRY A. MARZULLO, JR.
REGISTRATION NUMBER: 20,910
REFERENCE/DOCKET NUMBER: P-13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (914) 723-4300
   GENERAL INFORMATION:
  TELEFAX: (914) 723-4301
NFORMATION FOR SEQ ID NO: 14:
   MOLECULE TYPE: pe
HYPOTHETICAL: NO
FRAGMENT TYPE: in
ORIGINAL SOURCE:
  SOFTWARE: Word Perfect 5.0 for Windows 95.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/930,917A
FILING DATE: 16-Sep-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of COMPUTER: Compatible OPERATING SYSTEM: Will CONTROL OF CONTROL 
   APPLICANT:
   CORRESPONDENCE ADDRESS:
  TITLE OF INVENTION:
  APPLICANT:
  STREET:
   ORGANISM: Hur
STRAIN: IIIB
  TOPOLOGY:
  OF SEQUENCES:
   10583
  4, Application US/08930917A 6146635
   Scarsdale
  New York
Y: U.S.
   RAFVTI 13
   RAFVTI 6
   amino acid
  13 amino acids
15 Amino acid residues
  One Chase Road
  QUINTANA V. D.
G MEZ RODR GUEZ, C. B.
SILVA RODR GUEZ, R. C.
NAZ BAL G LVEZ, C.
   LEAL ANGULO, M. J
  ALVAREZ ACOSTA,
CARPIO MU%OZ, E
  GUILL N NIETO,
  DUARTE CANO,
  MART N DUNN,
   E: Floppy disk 3.5'' (1.4 MB).
Compatible PC IBM (80486, 8 M Ram).
   linear
   Human Immunodeficiency Virus Type
   Lackenbach Siegel Marzullo Aronson & Greenspar
  internal
   peptide
   Windows 95
   71.8%; Score 28; DB 100.0%; Pred. No. 14 tive 0; Mismatches
   System for the expression of heterologous
   /label= peptide
/note= "Active peptide of HIV-I envelope
from strain IIIB"
  Ġ
  G.≱
  DB 2;
   proteins
  <u>.</u>
   Length 13;
  Indels
   ٥.
   Gaps
   0
  glycopr
   Ś
   US-08-930-917A-14
  US-08-493-235-23
   Matches
  Patent No.
   Sequence 23,
  Query Match
   FEATURE:
   APPLICANT:
  APPLICANT:
   APPLICANT:
   APPLICANT:
   TOPOLOGY:
  COUNTRY:
  TELEPHONE:
   ENGTH:
   ξ
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COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/493,235
FILING DATE: 20)June)1995
CLASSIFICATION: 530
ATTORNEY/ACPNT ****
  Best Local Similarity
   TELEFAX: 619-235-0176 INFORMATION FOR SEQ ID NO:
   GENERAL INFORMATION:
MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
FRAGMENT TYPE:
  HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: Internal fragment
ORIGINAL SOURCE:
ORGANISM: VIH-1
INDIVIDUAL ISOLATE: IIIB
   CORRESPONDENCE ADDRESS:
ADDRESSEE: KNOBBE, MARTENS, OLSON AND BEAR
STREET: 620 NEWPORT CENTER DRIVE 16TH FLOOR
  TITLE OF INVENTION: PEPTIDES FOR USE IN VACCINATION AND TITLE OF INVENTION: INDUCTION OF NEUTRALIZING ANTIBODIES AGAINST HUMAN TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS
   MOLECULE TYPE: Pe
   SEQUENCE CHARACTERISTICS
  TELECOMMUNICATION INFORMATION:
  OTHER INFORMATION:
OTHER INFORMATION:
  NAME: Kaiser, AnneMarie REGISTRATION NUMBER: 37, REFERENCE/DOCKET NUMBER:
   CITY: NEWPORT BEACH
  STRANDEDNESS:
  STRANDEDNESS:
   10 RAFVTI
   1 RAFVTI 6
  6; Conservative
   3, Application US/08493235
5840313
  : 25 amino acids amino acid
   Amino acid
   USA
  Rymo, Lars
  Svennerholm, Bo
  Horal, Peter
  Jeansson, Stig
   Vahlne, Anders
  linear
  Unknown
                            NO peptide
  Peptide
  Unknown
  single
  100.0%;
   71.8%;
  41
   Central region of the loop V3 belonging to the protein gp120 from the VIH-1, isolation IIIB.
  37,649
   23:
   METRICS.12CPC1
  <u>.</u>.
  Score 28;
Pred. No.
  Mismatches
  DB 3;
16;
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   Length 15;
  Indels
  0;
  Gaps
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Matches

Conservative

0

Mismatches

0

Gaps

0

Query Match Best Local Similarity

71.8%; Score 28; 100.0%; Pred. No.

DB 26;

2

Length 25; ); Indels

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US-08-279-906A-26
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GENERAL INFORMATION:
   Sequence 11814, Application US/09902540
  Query Match
  Patent No. 561892
     APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
APPLICANT: Wiegand, Roger C.
APPLICANT: Myxococcus xanthus Genome Sequences and Uses Thereof
PILE REFERENCE: 38-10 (15849)8
   atent No. Joe
GENERAL INFORMATION:
Ohno, Tsuneya
CURRENT APPLICATION NUMBER: US/09/902,540
  INFORMATION FOR SEQ ID NO:
   SEQUENCE CHARACTERISTICS:
   ATTORNEY/AGENT INFORMATION:
NAME: No. 5618922and, Greta
REGISTRATION NUMBER: 35,302
  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/279,906A
FILING DATE:
CLASSIFICATION: 424
CLASSIFICATION: 424
   APPLICANT: Terada, Masaki
APPLICANT: Yoneda, Yukio
TITLE OF INVENTION: NM03 Antibody Materials and Methods
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
  MOLECULE TYPE:
   REFERENCE/DOCKET NUMBER: 32
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
  TELEPAX: 151-
TELEPAX: 25-3856
   STREET: 65.
STREET: Chicago
   Local Similarity
  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
  TOPOLOGY:
  LENGTH:
   COUNTRY:
  ADDRESSEE: Marshall, O'Toole, Gerstein, Murray
   20 RAFVTI 25
  σ
  f: 20 amino acids
amino acid
   60606
   RAFVTIGK 8
  RAFVTI 6
   Illinois
   Application US/08279906A
  6300 Sears Tower, 233 S. Wacker Drive
  (312) 474-0448
   USA
   Conservative
  linear
   peptide
  66.7%;
   Greta E.
   Score 26;
Pred. No.
  32028
  7:
   2
  Length 20,
   0,
   Gaps
   0
```

```
문
   δ
  RESULT 222
US-08-986-234-17
   밁

/ ORGANISM: Myxococcus xanthus
US-09-902-540-11814
  US-08-333-565-16
  US-08-986-234-17
  CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 11814
LENGTH: 23
   SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 17
LENGTH: 15
   GENERAL INFORMATION:
   Patent No. 5622852
   Sequence 16, Application US/08333565
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  Query Match
Best Local (
   Sequence 17, Appli
Patent No. 5981706
  Best Local Similarity
Matches 5; Conserv
  Matches
   Query Match
  CURRENT APPLICATION NUMBER: US/08/986,234
CURRENT FILING DATE: 1997-12-05
NUMBER OF SEQ ID NOS: 114
   APPLICANT: Wallen, et al.
TITLE OF INVENTION: Methods for Synthesizing Heat Shock Protein Complexes
FILE REPERENCE: UNME-0008-1
   GENERAL INFORMATION:
   ORGANISM: Human immunodeficiency virus
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   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM CC compatible
COMPUTER: Floppy disk
COMPUTER: IBM CC compatible
COMPUTER: IPM CC compatible
COMPUTER: POLDS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/333,565
FILING DATE: 31-OCT-1994
  APPLICANT: KORSMEYER, Stanley J.
TITLE OF INVENTION: Bcl-x/Bcl-2 ASSOCIATED CELL DEATH
TITLE OF INVENTION: REGULATOR
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
   APPLICANT:
  Local Similarity hes 5; Conserv
   COUNTRY:
   CITY: Palo Alto
   STREET:
   2 AFVTIGK 8
  94301
   σ
   RAFVTICK 8
   AFVTYGR 12
   California
  Application US/08986234
   379 Lytton Avenue
  Conservative
   Conservative
   31-OCT-1994
  64.1%;
  66.7%;
   Score 25; DB
Pred. No. 69;
1; Mismatches
  Score 26;
Pred. No.
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  4
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  Gaps
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  0
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δ
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   Ś
   US-08-333-565-16
  TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 16
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
STRANDEDNESS: single
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  REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15726
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2420
TELEPAX: (415) 326-2422
TELEPAX: (415) 00: 16:
  Patent No. 5834209
   APPLICATION NUMBER: US 08/333
FILING DATE: 31-OCT-1994
ATTORNEY/AGENT LIMPORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1572
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
   TELEFAX: (415) 326-24
INFORMATION FOR SEQ ID NO:
  GENERAL INFORMATION:
   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
ADDITICATION NUMBER: US 08/333,565
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LENGTH: 17 amino acids
TYPE: amino acid
STRANDEDNESS: single
  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 11-JUN-19
   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
  APPLICANT: KORSMEYER, Stanley J.
TITLE OF INVENTION: Bcl-x/Bcl-2 ASSOCIATED CELL DEATH
TITLE OF INVENTION: REGULATOR
   MOLECULE TYPE:
   NUMBER OF SEQUENCES: 59
  y Match 64.1%;
Local Similarity 66.7%;
les 4; Conservative
   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
  TOPOLOGY: linear
   SOFTWARE: PatentIn Release #1.0, Version #1.25
  COUNTRY:
  STREET: 379 Lytton Avenue CITY: Palo Alto
  TOPOLOGY:
   DDRESSEE:
3 FVTIGK 8
  12 FMTLGK 17
  3 FVTIGK 8
  94301
   4; Conservative
   California
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  linear
   Townsend and Townsend Khourie and Crew
  11-JUN-1995
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66.7%;
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   15726A-000700
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  Gaps
  Gaps
  0,
  0
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Query Match
Best Local Similarity
Tatches 5; Conserve
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   밁
  US-08-323-192D-51
   US-08-100-118-6
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US-08-100-118-6
  COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/100,118
FILING DATE: 19930730
CLASSIFICATION: 435
ATTORNEY/ACENT - 435
   CLONE: Consensus Sequence (CS) of HIV-1 V3 loop Patent No. 5580773
  Sequence 51, Application US/08323192D Patent No. 5786199
   Sequence 6, Application US/08100118 Patent No. 5580773
   Patent No.
  GENERAL INFORMATION:
   TELEFAX: 612-332-9081
INFORMATION FOR SEQ ID NO:
   GENERAL INFORMATION:
  NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
   IMMEDIATE SOURCE:
   TOPOLOGY: 1i
   SEQUENCE CHARACTERISTICS:
LENGTH: 18 amino acide
   TELECOMMUNICATION INFORMATION: TELEPHONE: 612-332-5300
   TITLE OF INVENTION: Design, Construction and Expression TITLE OF INVENTION: of Chimeric Proteins for Development of AIDS TITLE OF INVENTION: Vaccines and Diagnostic Reagents
  TITLE OF INVENTION:
  APPLICANT: Palese, TITLE OF INVENTION:
  NUMBER OF SEQUENCES:
  CORRESPONDENCE ADDRESS:
   APPLICANT:
                                   CITY: New York STATE: New Yor
  NAME: Strodthoff, Kristine M. REGISTRATION NUMBER: 34,259
REFERENCE/DOCKET NUMBER: 8682
  COUNTRY: U
ZIP: 55402
   STREET: 3100 No. 5
CITY: Minneapolis
                 COUNTRY:
  STREET:
   TYPE: amino acid
  ADDRESSEE:
  ADDRESSEE:
  11 RAFYTTGE 18
  1 RAFVTIGK 8
'RY: USA
10036-2711
                                New York
  逶
  1155 Avenue of the Americas
  E: Merchant & Gould
3100 No. 5580773west Center
   18 amino acids
  Conservative
  Kang, Lizhong
  USA
   linear
  Pennie & Edmonds
   peptide
   Chil-Yong
  EXPRESSION SYSTEMS AND VACCINES
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  RECOMBINANT NEGATIVE STRAND RNA VIRUS
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   8682.6-US-01
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Pred. No. 82;
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RESULT 227
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/975,699
FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/619,512
APPLICATION NUMBER: US 08/619,512
APPLICATION NUMBER: PCT/JP95/01515
FILING DATE: 31-JUL-1995
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APPLICATION NUMBER: PCT/JP95/01515
FILING DATE: 31-JUL-1995
PRIOR APPLICATION DATA:
   Sequence 15,
  TELEFAX: (212)869-974
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
  GENERAL INFORMATION:
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   SOPTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/323,192D FILING DATE: 14-OCT-1994
CLASSIFICATION: 435
   SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
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  TOPOLOGY: unknown
MOLECULE TYPE: peptide
  TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 790-9090
   ATTORNEY/AGENT INFORMATION:
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
  CORRESPONDENCE ADDRESS:
   NUMBER OF SEQUENCES:
   APPLICANT: TASAKA, HIROMICHI
TITLE OF INVENTION: ANTI-AIDS SECRETORY RECOMBINANT BCG
TITLE OF INVENTION: VACCINE
  APPLICANT:
  APPLICANT:
   PPLICANT:
   STREET: 1755 S.
CITY: ARLINGTON
   NAME: Coruzzi, Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7682-035
   MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
   STRANDEDNESS:
   COUNTRY: U
  TELEPHONE:
   ADDRESSEE: OBLON. SPIVAK, MCCCLELLAND, MAIER & NEUSTADT.
ADDRESSEE: P.C.
   10 RAFYTTGE 17
  1 RAFVTIGK 8
  amino acid
   Ş
   Application US/08975699
  E: P.C.
1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
  USA
   (212)869-9741/8864
  YAMAZAKI, AKIHIRO
   YAMAKAZI,
  HONDA, MITSUO
  MATSUO,
   YOSHITOMO
JP 178462/1994
   1; Mismatches
  Score 25; DB
Pred. No. 87;
  DB 1;
   2; Indels
  Length 19;
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```

```
맑
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   US-08-972-089-15
  Sequence 15,
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  Query Match
   Patent No.
           APPLICATION NUMBER: PCT/JP95/01515
PRILING DATE: 31-JUL-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 178462/1994
PILING DATE: 29-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-795-0X PCT
   TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO:
  GENERAL INFORMATION:
   FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION UDMBER:
APPLICATION NUMBER:
FILING DATE:
  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/972,089
  ZIP: 22202
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
  MOLECULE TYPE: peptide ORIGINAL SOURCE:
  SEQUENCE CHARACTERISTICS:
   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
   APPLICANT: TASAKA, HIROMICHI
TITLE OF INVENTION: ANTI-AIDS SECRETORY RECOMBINANT BCG
TITLE OF INVENTION: VACCINE
   TELECOMMUNICATION INFORMATION: TELEPHONE: 703-413-3000
  APPLICANT:
   ATTORNEY/AGENT INFORMATION:
   APPLICANT:
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   APPLICANT:
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TELECOMMUNICATION INFORMATION
   64.1%;
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les 5; Conservative
   COUNTRY:
   STREET: 1755 S. CITY: ARLINGTON
  REGISTRATION NUMBER: 24
REFERENCE/DOCKET NUMBER:
   LENGTH: 19 amino acids
TYPE: amino acid
STRANDEDNESS: single
  NAME: OBLON,
  FILING DATE:
  ADDRESSEE: P.C.
STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
  TOPOLOGY: linear
  ADDRESSEE:
  TELEPHONE:
  12 RAFYTTGE 19
  1 RAFVTICK 8
  Ş
  Application US/08972089
   USA
  703-413-2220
  HONDA, MITSUO
  MATSUO, KAZUHIRO
CHUJO, YOSHITOMO
   YAMAKAZI,
   YAMAZAKI, AKIHIRO
   HUMAN IMMUNODEFICIENY VIRUS
  OBLON. SPIVAK, MCCCLELLAND, MAIER & NEUSTADT,
  Floppy disk
  NORMAN F
  SHUDO
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   24,618
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  Mismatches
  2
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LENGTH: 19 amino acids
  REGISTRATION NUMBER: 26,154
REFERENCE/DOCKET NUMBER: 29
TELECOMMUNICATION INFORMATION:
   FILING DATE: 28-FEBRUARY-1990 (AHANDONED)
APPLICATION NUMBER: USSN 07/687,376
FILING DATE: 18-APRIL-1991 (ABANDONED)
APPLICATION NUMBER: USSN 08/327,636
FILING DATE: 24-OCTOBER-1994 (ABANDONED)
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Richard S
NAME: Clark, Richard S
  SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION WHEER: US/08/363,276B
FILLING DATE: 22-DECEMBER-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 07/486,546
                  MOLECULE TYPE:
ORIGINAL SOURCE:
   COMPUTER READABLE FORM:
   TITLE OF INVENTION:
   MOLECULE TYPE: peptide ORIGINAL SOURCE:
   CORRESPONDENCE ADDRESS
  NUMBER OF SEQUENCES:
  APPLICANT:
   ORGANISM:
  STRANDEDNESS:
   TELEFAX: 212-765-2519
  TELEPHONE:
   COMPUTER: IBM CON OPERATING SYSTEM:
  TOPOLOGY:
  MEDIUM TYPE:
  COUNTRY: USA
ZIP: 10112-0228
   STREET: 30 Roc
CITY: New York
  Local Similarity
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   ORGANISM: HUM
STRAIN: HIV-1
   TYPE: amino acids
   ADDRESSEE:
  STRANDEDNESS:
  TELEPHONE:
  COPOLOGY:
   12 RAFYTTGE 19
  1 RAFVTIGK 8
  amino acid
  Y
  B: Raymond
30 Rockefeller Plaza
   ಟರ್ಟ single
linear
   linear
Human Immunodefficiency Virus Type 1
   BONA ET AL
  Conservative
  HUMAN IMMUNODEFICIENY VIRUS
  IBM Compatible
   Brumbaugh, Graves,
  703-413-3000
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COMPRISING ANTIGEN
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   15:
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  2;
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5
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  Sequence 1, Applic
Patent No. 6204250
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  Query Match
  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
  GENERAL INFORMATION:
   NAME: Clark, Richard S
REGISTRATION NUMBER: 26,154
REFERENCE/DOCKET NUMBER: 291
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-408-2558
  FEATURE:
   MOLECULE TYPE: peptide ORIGINAL SOURCE:
  FILING DATE:
ATTORNEY/AGENT INFORMATION:
  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
  COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
  ZIP: 10112-0228
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
  APPLICANT: BOT and E
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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   LOCATION: 301...319
OTHER INFORMATION: Envelope Protein gp120
   NAME/KEY:
   ORGANISM:
  APPLICATION NUMBER: FILING DATE: 22-NO
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  TELEFAX: 212-765-2519
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  ADDRESSEE: Brumbaugh, Graves, Donohue & ADDRESSEE: Raymond
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   COUNTRY:
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   10 RAFYTTGE 17
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   Application US/08755034
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   22-NOVEMBER-1996
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87;
  ب
   Length 19;
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10 RAFYTTGE 17

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  US-10-125-594-6
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Patent No. 6740747
  Query Match
   SOFTWARE: PatentIn version 3.1

SEQ ID NO 6

LENGTH: 19
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PRIOR FILING DATE: 2001-04-20
NUMBER OF SEQ ID NOS: 11
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
  FILE REFERENCE: 12837-4
CURRENT APPLICATION NUMBER: US/10/125,594
CURRENT FILING DATE: 2002-04-19
   ENERAL INFORMATION:
  APPLICANT: Kaushik, Azad
APPLICANT: Saini, Surin
   ATTORNEY AGENT INFORMATION:
NAME: Clark, Richard S
REGISTRATION NUMBER: 26,154
REFERENCE/DOCKET NUMBER: 2988
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-408-2558
  SENERAL INFORMATION:
  RPPLICANT: Saini, Surinder Singh
RITLE OF INVENTION: No. 6740747el Bovine VDJ Cassette, BF1H1, Suitable for Antigeniza
   quence 1, Application PC/TUS9516718
  TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
   OTHER INFORMATION: Human HIV-1
   APPLICANT: MOUNT SINAI SCHOOL OF MEDICINE OF THE APPLICANT: CITY UNIVERSITY OF NEW YORK TITLE OF INVENTION: CHIMERIC ANTIBODIES
   SOFTWARE: FASTSEQ VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/
   COMPUTER READABLE FORM:
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APPLICATION NUMBER:
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Local Similarity 62.5%;
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   STREET: 30 Rocl
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  1 RAFVTIGK 8
  10112-0228
  30 Rockefeller Plaza
  USA
   212-765-2519
  IBM Compatible
  Raymond
   Diskette
  COMPRISING ANTIGEN BINDING SITES AND B AND T CELL EPITOPES
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  29889-165/29528
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Best Local Similarity
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOPTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/08:
  INFORMATION FOR SEQ ID NO:
   CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: 08/477
APPLICATION UNMBER: 1995
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Richard S
REGISTRATION NUMBER: 26,15
  MOLECULE TYPE:
ORIGINAL SOURCE
  SEQUENCE CHARACTERISTICS
LENGTH: 19 amino acid
   MOLECULE TYPE:
ORIGINAL SOURCE:
  REFERENCE/DOCKET NUMBER: 29
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-408-2558
  APPLICANT: UNIVERSI
  FEATURE:
   CORRESPONDENCE ADDRESS:
  NUMBER OF SEQUENCES:
  APPLICANT: MOUNT SINAI SCHOOL OF MEDICINE OF THE CITY
   LOCATION: 301...319
OTHER INFORMATION:
   ORGANISM: Human Immunodefficiency Virus Type
                                   NAME/KEY:
  TYPE: amino o
   ORGANISM:
   TOPOLOGY:
  TELEFAX:
  CITY: New York
   STREET:
   NAMB/KBY:
  STRANDEDNESS:
  LENGTH:
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   10 RAFYTTGE 17
   1 RAFVTIGK 8
  υ
••
   amino acids
   amino acid
  Application PC/TUS9608995
   E: Brumbaugh, Graves, Donohue & Raymond
30 Rockefeller Plaza
  19 amino acids
  UNIVERSITY OF NEW
  Conservative
  212-765-2519
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   Human Immunodefficiency Virus Type 1
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  PEGYLATED MODIFIED PROTEINS
   64.1%;
Envelope Protein gp120
   08/477,421
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Query Match

Score 25;

DB 5;

Length 19;

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   Best Local Similarity 62.5%; Matches 5; Conservative
   GENERAL INFORMATION:
APPLICANT: Clark, Ross Gl
APPLICANT: Lownan, Henry B.
APPLICANT: Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
   TELEFAX: (312) 474-044
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
  GENERAL INFORMATION:
APPLICANT: Ohno, Tsuneya
   ATTORNEY/AGENT INFORMATION:
NAME: NO. 5618922and, Greta E
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 3202
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
   TOPOLOGY: li
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
  STREET: Chicago
CITY: Chicago
Illinois
INSA
                              CORRESPONDENCE ADDRESS:
  NUMBER OF SEQUENCES:
   SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
   CORRESPONDENCE ADDRESS:
  APPLICANT: Terada, Masaki
APPLICANT: Yoneda, Yukio
TITLE OF INVENTION: NM03 Antibody Materials and Methods
NUMBER OF SEQUENCES: 27
   Local Similarity 62.5%; les 5; Conservative
 ADDRESSEE:
STREET: 1
   TYPE: amino acid
  TELEPHONE: (312) 474-0448
  FILING DATE:
CLASSIFICATION:
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   DDRESSEE:
  ADDRESSEE: Marshall, O'Toole, Gerstein, Murray &
   10 RAFYTTGE 17
  1 RAFVTIGK 8
  60606
   RAFVTIGK 8
   Application US/08825852
  Application US/08279906A
1 DNA Way
  6300 Sears Tower, 233 S. Wacker Drive
   20 amino acids
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   424
  US/08/279,906A
  Greta E.
   1; Mismatches
  Score 25; DB
Pred. No. 91;
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   Pred. No. 87;
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TELEPHONE: 650/225-1896
TELEPAX: 650/952-9881
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LENGTH: 20 amino acids
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  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 20 amino acide
   ATTORNEY/AGENT INFORMATION:
NAME: HASAK, JANET E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
   NUMBER OF SEQUENCES: 7
  ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,
  OPERATING SYSTEM: PC-DOS/MS-DC SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
  APPLICANT: Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
   APPLICANT: Lowman, Henry B.
APPLICANT: Lowman, Henry B.
   TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-1896
   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inc
   FILING DATE: 04-Ap
CLASSIFICATION: 51
   COUNTRY: U
   STREET: 1 DNA Way
CITY: South San Francisco
   LENGTH: 20 amino acids TYPE: Amino Acid
  SOFTWARE: WinPatin
   APPLICATION NUMBER: FILING DATE: 04-Apr CLASSIFICATION: 514
   CITY: South San Francisco
STATE: California
   ADDRESSEE:
   REFERENCE/DOCKET NUMBER:
   MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible
  14 ÁFMAVGK 20
   2 AFVTIGK 8
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  94080
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   Conservative
  USA
  650/952-9881
   Linear
   Genentech, Inc.
   04-Apr-1997
   04-Apr-1997
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  (Genentech)
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   US/08/825,852
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                GENERAL INFORMATION:
APPLICANT: Clark, Ross Gl
APPLICANT: Lowman, Henry B.
APPLICANT: Robinson, Iain C.A.F.
APPLICANT: Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: Amino Acid
   ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P107
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
  GENERAL INFORMATION:
  APPLICANT: Clark, Nous B.
APPLICANT: Lowman, Henry B.
APPLICANT: Robinson, Tain C.A.F.
APPLICANT: Robinson, Islin C.A.F.
APPLICANT: Nous Tain C.A.F.
APPLICANT: Lowman, Henry B.
APPLICANT: Robinson, Island C.A.F.
  MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION NUMBER: US/09/052,888 FILING DATE: 31-Mar-1998 CLASSIFICATION: 514
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MEDIUM TYPE: 3.5 inc
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CITY: South San Francisco
STATE: California
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  TOPOLOGY:
  TELEFAX:
   DDRESSEE:
   14 AFMAVGK 20
  14 AFMAVGK 20
  2 AFVTICK 8
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   94080
   AFVTIGK 8
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   Application US/09052888
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57.1%;
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  1; Indels
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   Length 20;
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  US-09-052-888-65
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/052,88
EILING DATE: 31-Mar-1998
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1071P1
TELECOMMUNICATION INFORMATION:
   TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
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LENGTH: 20 amino acids
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
  TYPE: Amino Acid
   TOPOLOGY:
   TELEPHONE:
   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/723,890
FILING DATE: 28-No. 6608031-2000
CLASSIFICATION: 514
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1071P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/052,888
APILING DATE: 31-Mar-1998
ATTORNEY/AGENT INFORMATION:
  COMPUTER READABLE FORM:
  CORRESPONDENCE ADDRESS:
  Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
   APPLICANT: Clark, Ross Gl
   14 AFMAVGK 20
   2 AFVTICK 8
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   4
   Similarity
   South San
   STATE: California
   MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
   ZIP: 94080
  COUNTRY: USA
  CITY: South San
  California
  ADDRESSEE: Genentech, Inc.
  Application US/09723890
  Conservative
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  650/225-1896
  Jowman, Henry B.
   1 DNA Way
   Francisco
   64.1%;
57.1%;
  US/09/052,888
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  멍
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Best Local :
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Patent No. 60
  Query Match
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
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APPLICATION NUMBER: US/09/723,890
FILING DATE: 28-No. 6608031-2000
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
  APPLICATION NUMBER: US/09/052,888 FILING DATE: 31-Mar-1998 ATTORNEY/ACENT INFORMATION:
   REFERENCE/DOCKET NUMBER: P1071P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
  Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
   TYPE: Amino Acid
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 64:
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  CORRESPONDENCE ADDRESS
  APPLICANT: Clark, Ross Gl
  14 AFMAVGK 20
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  4; Conservative
  Similarity
   AFVTIGK 8
  CITY: South San F. STATE: California
   NAME: Hasak, Janet E. REGISTRATION NUMBER: 28,616
  TOPOLOGY: Linear
   TYPE: Amino Acid
   TELEFAX: 650/952-9881
  COUNTRY: USA
   STREET:
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 Application US/09723901
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  Application US/09723890
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   1 DNA Way
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GENERAL INFORMATION:
APPLICANT: Clark, Ross Gl
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GENERAL INFORMATION:
APPLICANT: Clark, Ross Gl
  NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1071P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
TELEPHONE: 650/25-1896
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/723,901
FILING DATE: 28-No. 6620789-2000
  CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA WAY
  Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
   APPLICATION NUMBER: 09/052,888 FILING DATE: 31-Mar-1998 ATTORNEY/AGENT INFORMATION:
  ZIP: 94080
COMPUTER READABLE FORM:
   PRIOR APPLICATION DATA:
  NUMBER OF SEQUENCES: 109
CORRESPONDENCE ADDRESS:
   Lowman, Henry B.
Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
  14 AFMAVGK 20
  2 AFVTICK 8
   4; Conservative
  CITY: South San Francisco
STATE: California
  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
   CITY: South San F:
STATE: California
COUNTRY: USA
   ZIP: 94080
   TYPE: Amino Acid
  COUNTRY: USA
   CLASSIFICATION: <Unknown>
   ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
   MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
  Application US/09723901
   ENGTH: 20 amino acids
   64.1%;
57.1%;
  Francisco
   <u>ب</u>
   Score 25; DB
Pred. No. 91;
  Mismatches
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  Length 20
   Indels
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,
   Gaps
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APPLICATION NUMBER: US/09/723,901
PILING DATE: 28-No. 6620789-2000
CLASSIFICATION: UNKNOWN>
PRIOR APPLICATION DATA:

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SEQUENCE DESCRIPTION: US-09-723-547-64
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  US-09-723-901-65
   Sequence 64, Applicati
Patent No. 6632794
GENERAL INFORMATION:
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  INFORMATION FOR SEQ ID NO: 64: SEQUENCE CHARACTERISTICS:
   INFORMATION FOR SEQ ID NO: 65:
  PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/052,888

PILING DATE: «Unknown»

ATTORNEY/AGENT INFORMATION:

NAME: Hasak, Janet E.

REGISTRATION NUMBER: 28,616

REGISTRATION NUMBER: P1071P1

TELECOMMUNICATION INFORMATION:

TELEPHONE: 650/225-1896
   ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/723,547
FILING DATE: 28-No. 6632794-2000
CLASSIFICATION: <Unknown>
  NAME: HaBAK, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
TELEFAX: 650/952-9881
  Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
  TOPOLOGY: Linear SEQUENCE DESCRIPTION: SEQ ID NO: 65:
   SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: Amino Acid
   APPLICATION NUMBER: 09/052,888
FILING DATE: 31-Mar-1998
ATTORNEY/AGENT INFORMATION:
  CORRESPONDENCE ADDRESS:
   APPLICANT: Clark, Ross Gl
   14 AFMAVGK 20
   2 AFVTIGK 8
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   TYPE: Amino Acid
  STATE: California
  TELEFAX: 650/952-9881
   COUNTRY: USA
  CITY: South San
   ADDRESSEE: Genentech, Inc
STREET: 1 DNA Way
  ENGTH: 20 amino acids
   ADDRESSEE:
  Application US/09723547
  Conservative
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57.1%;
  Francisco
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Patent No. 6635619
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Best Local Similarity
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   Patent No. 6632794
GENERAL INFORMATION:
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   Best Local Similarity
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   TELEFAX: 650/952-9881
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Windatin (Genencech)
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FILING DATE: 28-No. 6632794-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
  APPLICATION NUMBER: 09/052,888 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
   ROBINSON, Iain C.A.F. TITLE OF INVENTION: Insulin-like NUMBER OF SEQUENCES: 109
   REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
  SEQUENCE DESCRIPTION: SEQ ID NO: 65:
  CORRESPONDENCE ADDRESS
  14
  14 AFMAVGK 20
   2 AFVTIGK 8
  2 AFVTIGK 8
   4; Conservative
  AFMAVGK 20
  TYPE: Amino Acid
TOPOLOGY: Linear
  NAME: Hasak, Janet E. REGISTRATION NUMBER: 28,616
  STATE: California
  COUNTRY: USA
  CITY: South San Francisco
  Application US/09724127
  ENGTH: 20 amino acids
   TELEPHONE: 650/225-1896
   ZIP: 94080
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  Application US/09723547
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57.1%;
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57.1%;
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Pred. No. 91;
   Score 25; DB
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   Gaps
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Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109

Lowman, Henry B

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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/724,127
FILING DATE: 28.No. 6635619-2000
CLASSIFICATION. CUNKNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/052,888 FILING DATE: 31-Mar-1998 ATTORNEY/AGENT INFORMATION:
  Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
   APPLICATION NUMBER: 09/052,888
FILING DATE: 31-Mar-1998
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
   CURRENT APPLICATION DATA:
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FILING DATE: 28-No. 6635619-2000
CLASSIFICATION: <Unknown>
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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   PRIOR APPLICATION DATA:
  CORRESPONDENCE ADDRESS:
   14 AFMAVGK 20
  2 AFVTIGK 8
  4; Conservative
   CITY: South San Francisco
STATE: California
   ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
  ZIP: 94080
   COUNTRY:
  TYPE: Amino Acid
  TELEFAX: 650/952-9881
  ADDRESSEE: Genentech, Inc
STREET: 1 DNA Way
   STATE: California
  CITY: South San
   Lowman, Henry B.
   USA
   WinPatin (Genentech)
  64.1%;
  Francisco
  Mismatches
   Score 25; DB Pred. No. 91;
   4.
  1; Indels
  Length 20;
  Gaps
  0,
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밁
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  US-09-723-931-64
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  Sequence 64, Application Patent No. 6645775
GENERAL INFORMATION:
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  Matches
   Query Match
Best Local Similarity
   TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
  INFORMATION FOR SEQ ID NO: 65:
   COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb i
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
  NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,6
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
  APPLICATION NUMBER: 09/052,888-FILING DATE: 31-Mar-1998
ATTORNEY/AGENT INFORMATION:
  CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
   SEQUENCE DESCRIPTION: SEQ ID NO: 64:
   PRIOR APPLICATION DATA:
   Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
   SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: Amino Acid
   TOPOLOGY: Linear SEQUENCE DESCRIPTION:
   REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
   APPLICANT: Clark, Ross Gl
2 AFVTIGK 8
  14 AFMAVGK 20
  Similarity
4; Conserva
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  4; Conservative
  APPLICATION NUMBER: US/09/723,931 FILING DATE: 28-No. 6645775-2000 CLASSIFICATION: <Unknown>
   LENGTH: 20 amino acids TYPE: Amino Acid
   STATE: California COUNTRY: USA
   CITY: South San
  Application US/09723931
   TELEFAX: 650/952-9881
  TELEPHONE: 650/225-1896
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  94080
  owman, Henry B.
   64.1%;
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57.1%;
   Francisco
   inch, 1.44 Mb floppy disk
   SEQ ID NO: 65:
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Pred. No. 91;
2; Mismatches
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   Score 25; DB Pred. No. 91;
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  DB 4; Length 20
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14 AFMAVGK 20

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RESULT 248
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; Sequence 65, Application
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US-09-723-873-64
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Patent No. 6677305
Patent No. 10677301
GENERAL INFORMATION:
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  CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/723,931
FILING DATE: 28-No. 6645775-2000
CLASSIFICATION: <Unknown>
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/052,888
FILING DATE: 31-Mar-1998
ATTORNEY/AGENT INFORMATION:
   LOWMAN, Henry B.
Robinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
   TOPOLOGY: Linear SEQUENCE DESCRIPTION: SEQ ID NO: 65:
  NAME: HASAK, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
  RObinson, Iain C.A.F.
TITLE OF INVENTION: Insulin-like Growth Factor Agonist Molecules
NUMBER OF SEQUENCES: 109
   APPLICANT: Clark, Ross Gl
  APPLICANT: Clark, Ross Gl
  CORRESPONDENCE ADDRESS:
   CORRESPONDENCE ADDRESS
   14 AFMAVGK 20
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   4; Conservative
ADDRESSEE: Genentech, Inc. STREET: 1 DNA Way CITY: South San Francisco STATE: California COUNTRY: USA
  TYPE: Amino Acid
  STATE: California
  CITY: South San Francisco
  ADDRESSEE: Genentech, Inc
STREET: 1 DNA Way
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   Application US/09723931
   94080
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  64.1%;
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  P1071P1
  DB 4; Length 20;
   1; Indels
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US-09-723-873-65
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TELEFAX: 650/952-9881
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
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PILING DATE: 28-No. 6677305-2000
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PILING DATE: 31-Mar-1998
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FILING DATE: 31-Mar-1998
ATTORNEY/AGENT INFORMATION:
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TITLE OF INVENTION: Insulin-like
NUMBER OF SEQUENCES: 109
CORRESPONDENCE ADDRESS:
  NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
   TYPE: Amino Acid
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 64:
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   APPLICANT: Clark, Ross Gl
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   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
  14 AFMAVGK 20
   2 AFVTIGK 8
   4; Conservative
   NAME: Hasak, Janet E.
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REGISTRATION NUMBER: P1071P1
   STATE: California
  ZIP: 94080
   CITY: South San Francisco
   STREET:
   ADDRESSEE:
   LENGTH: 20 amino acids
CHARACTERISTICS:
   Lowman, Henry B.
  1 DNA Way
  64.18;
57.18;
  Genentech, Inc.
   2; Mismatches
  Score 25; DB
Pred. No. 91;
   P1071P1
  Growth Factor Agonist Molecules
  DB 4;
  Length 20;
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   Gaps
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TYPE: Amino Acid
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TOPOLOGY: Linear
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SEQUENCE DESCRIPTION: SEQ ID NO: 65:
US-09-723-873-65

Query Match
Best Local Similarity 57.1%; Score 25; DB 4; Length 20;
Best Local Similarity 57.1%; Pred. No. 91;
Matches 4; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
Qy 2 AFVTIGK 8

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Db 14 AFMAVGK 20

Search completed: May 16, 2005, 10:03:40

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